

Interlaboratory prevalidation of a new in vitro transcriptional activation assay for the screening of (anti-)androgenic activity of chemicals using the UALH-hAR cell line

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1	Interlaboratory prevalidation of a new in vitro transcriptional activation assay for the
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24	Keywords: androgen receptor; transactivation assay; stable luciferase reporter cell line;
25	interlaboratory study.

26 Abstract

27 We report an interlaboratory evaluation of a recently developed androgen receptor (AR) transactivation assay using the UALH-hAR reporter cell line that stably expresses the luciferase 28 gene under the transcriptional control of androgen receptor elements (AREs) with no 29 30 glucocorticoid receptor (GR) crosstalk. Herein, a two-step prevalidation study involving three laboratories was conducted to assess performance criteria of the method such as transferability 31 as well as robustness, sensitivity, and specificity. The first step consisted in the validation of 32 33 the transfer of the cell line to participant laboratories through the testing of three reference chemicals: the AR agonist dihydrotestosterone, the AR antagonist hydroxyflutamide and the 34 35 glucocorticoid dexamethasone. Secondly, a blinded study was conducted by screening a 36 selection of ten chemicals, including four AR agonists, five AR antagonists, and one non-active chemical. All test compounds yielded the same activity profiles in all laboratories. The logEC₅₀ 37 (agonist assay) or $logIC_{50}$ (antagonist assay) were in the same range, with intra-laboratory 38 coefficients of variation (CVs) of 0.1-3.4% and interlaboratory CVs of 1-4%, indicating very 39 good within- and between-laboratory reproducibility. Our results were consistent with literature 40 and regulatory data (OECD TG458). Overall, this interlaboratory study demonstrated that the 41 42 UALH-hAR assay is transferable, produces reliable, accurate and specific (anti)androgenic 43 activity of chemicals, and can be considered for further regulatory validation.

45

1. Introduction

Androgens are steroid hormones that play pivotal roles in the development and reproduction of 46 vertebrates, notably in the male reproductive system. Most of their biological effects are 47 mediated by their action on nuclear androgen receptor (AR), a ligand-dependent transcriptional 48 factor that regulates many downstream androgen-dependent signalling pathways. Disrupting 49 androgen signalling alters normal reproductive functions and can also account for several 50 pathophysiological functions, such as prostate cancer. The AR signalling pathway is for 51 instance an important target of anti-cancer drugs, with an active search for new AR inhibitors 52 in the context of prostate cancer therapies (Nigro et al., 2021). It is also now well documented 53 54 that AR signalling can be disrupted by chemicals exerting either androgenic or antiandrogenic 55 activities (Gray et al., 2001; Scott et al., 2009). Environmental chemicals such as pharmaceuticals (Runnalls et al., 2010), pesticides (Kojima et al., 2004; Lemaire et al., 2004; 56 Raun Andersen et al., 2002), or flame retardants (Hamers et al., n.d.) that can be found in air, 57 surface waters or food (Biedermann and Grob, 1998; Creusot et al., 2014; Haith and Rossi, 58 2003; Rodriguez-Mozaz et al., 2004; Russo et al., 2019; Sumpter, 2005), might pose risks to 59 wildlife and humans through the disruption of the AR-signalling pathway. Therefore, there is a 60 need for reliable tools to assess the (anti)androgenic activities of chemicals and their mixtures. 61 62 As such, several established stable reporter cell lines such as PALM (Molina-Molina et al., 2013, 2006; Térouanne et al., 2000), HELN-AR (ERα-DBD) (Delfosse et al., 2012; Grimaldi 63 et al., 2019), or MDA-kb2 (Aït-Aïssa et al., 2010; Wilson et al., 2002) or AR-EcoScreen (Satoh 64 et al., 2004) have been developed and successfully used to screen (anti)-androgenic activities 65 66 of chemicals or environmental mixtures. However, these reporter gene assays present a limit regarding AR-specificity of their response because of the use of promoter constructs that 67 possess response elements to other endogenously expressed nuclear receptors such the 68 69 glucocorticoid (GR) and progesterone (PR) receptors. Indeed, GR is naturally expressed in

many cell lines and, once activated by its ligands, has been shown to activate (PALM, MDA-70 71 kb2, AR-Ecoscreen) or repress (HELN-AR-(ERa-DBD)) AR-responsive luciferase constructs 72 in these AR-reporter cell lines. To overcome this limit, other in vitro cell lines have been developed for more specific assessment of androgenic activity of chemicals, either by knocking 73 74 out endogenous GR expression, such as in the AR-EcoScreen GR KO (Zwart et al., 2017) or 75 22Rv1/MMTV GR-KO (Park et al., 2021; Sun et al., 2016) cell lines, or by using a cell model that naturally expresses very low levels of GR, such as in the U2OS AR-CALUX assay 76 (Milcamps et al., 2021, 2020; Sonneveld et al., 2006, 2005; van der Burg et al., 2015, 2010). 77 All three methods are included in the OECD Guideline TG 458 (OECD, 2020). 78

To address the current needs of AR specific in vitro tools with as minimal GR-crosstalk as 79 possible, a new U2OS-derived luciferase reporter cell line, UALH-hAR, has been recently 80 developed (Dellal et al., 2020). This cell line expresses high levels of AR but not GR, as well 81 as the luciferase gene under the control of an ARE selective promoter (contains 6 AREs), which 82 83 makes the luciferase response very sensitive and selective to AR ligands in this cell model. Such model could represent a relevant tool for screening (anti)androgenic activity of test 84 substances notably within a regulatory context. In the present study, we conducted a two-step 85 86 interlaboratory exercise to assess the robustness of UALH-hAR cell line and test method as a potential transferable and reliable assay for the specific detection of androgenic and 87 antiandrogenic activity of chemicals, using a selection of ten substances, including AR agonists 88 (4) AR antagonists (5) and one inactive chemical. 89

90

2. Material and Methods

2.1.

91

Design of the interlaboratory prevalidation study

92 The prevalidation study involved three laboratories: laboratory A coordinated the 93 prevalidation, prepared and sent around test chemicals, collected, analysed and reported the

data (INERIS, Unité Ecotoxicologie des Substances et Milieux, UMR-I 02 SEBIO, Parc
ALATA, 60550 Verneuil-en-Halatte, France); laboratory B created the cell line, developed the
method and distributed the cell line (IRCM, Inserm U1194, 34290 Montpellier, France); finally
laboratory C served as a naive laboratory (Tame-Water, 85000 La Roche-sur-Yon, France). All
three laboratories performed the experiments as described thereafter.

The prevalidation was done in two phases. For phase I, each laboratory tested three reference 99 compounds in both agonist (exposure to test chemical alone) and antagonist (co-exposure to 100 test chemical with reference agonist dihydrotestosterone DHT 3.16×10⁻¹⁰ M) modes in three 101 102 independent runs, *i.e.*, separate experiments using cells from a different pool, freshly diluted chemicals, and conducted on different days. Selected compounds for the phase I were the 103 androgen receptor (AR) agonist dihydrotestosterone (DHT), the AR antagonist 104 hydroxyflutamide (HF) and dexamethasone (DEX), a synthetic GR agonist known to exert no 105 106 agonistic activity towards AR. These compounds were all sent as powders by the laboratory A 107 to laboratories B and C and each laboratory then proceeded to the preparation of stock solutions in dimethyl sulfoxide (DMSO). Results of each independent run and of each laboratory were 108 then compared qualitatively (*i.e.*, active or non-active compound) and quantitatively (based on 109 110 comparison of $logEC_{50}$ and $logIC_{50}$ obtained after modeling of concentration-response curves) to assess within laboratory reproducibility (WLR) and between laboratory reproducibility 111 (BLR). 112

For phase II, WLR and BLR were assessed in real testing conditions by screening a selection of ten chemical substances in a blind study in agonist and antagonist modes in three independent runs. Tested compounds included the three reference chemicals used in phase I, as well as three AR agonists ligands: testosterone (T), medroxyprogesterone acetate (MPA), and methyltrienolone (R1881); three AR antagonists: bisphenol A (BPA), bisphenol C (BPC), and vinclozolin M2 (M2V); and one compound with no expected AR effect: mono(2ethylhexyl)phthalate (MEHP), a metabolite of di(2-ethylhexyl)phthalate (DEHP). All stock
solutions in DMSO were prepared by laboratory A, then anonymized (coded) and sent to
laboratories B and C. At laboratory A, the anonymization of stock solutions was performed by
a person external to the study to ensure blind testing conditions in this lab. Sample codes were
different for each laboratory. Results were analysed as in phase I to evaluate the WLR and BLR
of the test method.

125 **2.2.** *Test chemicals*

Chemical structures and main properties of the ten selected substances are presented in Table 126 127 1. Dihydrotestosterone (DHT), hydroxyflutamide (HF), dexamethasone (DEX), methyltrienolone (R1881), testosterone (T), medroxyprogesterone acetate (MPA), bisphenol A 128 (BPA), bisphenol C (BPC) and mono(2-ethylhexyl)phthalate (MEHP) were obtained from 129 130 Sigma-Aldrich Chemical Co. (Saint-Quentin Fallavier, France), and vinclozolin M2 (M2V) was obtained from Santa Cruz Biotechnology, Inc. (Heidelberg, Germany). Stock solutions of 131 chemicals were prepared in dimethyl sulfoxide (DMSO) and stored at -20°C. Fresh dilutions of 132 test chemicals in cell culture medium were prepared before each run and the final DMSO 133 concentrations did not exceed 0.1% (v/v) or 0.2% (v/v) of the culture medium in agonist and 134 antagonist modes, respectively. All three laboratories used the same batch of powders (Phase 135 I) or stock solutions (Phase II), prepared and sent by the lead laboratory (Laboratory A). 136

137

2.3. The UALH-hAR cell line

The previously established UALH-hAR cell line (Dellal et al., 2020) was derived from human osteosarcoma cells U2OS using a two-step stable transfection with three plasmids (Fig. 1). To establish the UALH cell line, U2OS cells were first stably transfected with the 6xARE(RAD9)Luc+/Hygromycin plasmid containing the firefly luciferase gene under the control of six androgen response elements (AREs) from the AR and PR selective RAD9 gene

promoter and placed upstream of the collagenase promoter and containing the resistance gene 143 to hygromycin under the control of the SV40 promoter. Then, UALH cells were co-transfected 144 with (i) the expression vector pSG5-AR-Puromycin containing the gene coding for hAR under 145 the control of the simian virus promoter (SV40) and resistance gene to puromycin under the 146 147 control of the phosphoglycerate kinase promoter (PGK) and (ii) the expression vector pSG5-AR-Renilla-Neomycin containing the gene coding for hAR under the control of the SV40 148 promoter, the Renilla luciferase reporter gene under the control of the constitutive human 149 150 cytomegalovirus (CMV), the firefly luciferase gene under the control of six androgen response elements (ARE) from the RAD9 gene promoter and placed upstream of the collagenase 151 152 promoter, and the resistance gene to neomycin under the control of the SV40 promoter. The 153 third plasmid was included to overexpress both AR and luciferase expression. The AR sequence was chosen from the NCBI database under the ID NM_000044.3. Androgen responses elements 154 of from the RAD9 gene promoter were chosen due to their high AR selectivity versus GR 155 (Moehren et al., 2008). Cells were incubated in a 5% CO₂ humidified atmosphere at 37°C. 156 Selection of UALH-hAR cells was made with 0.5 µg/mL puromycin, 1 mg/L neomycin G418 157 and 0.25 mg/mL hygromycin. Cells are passaged after reaching 75-90% confluence and the 158 159 number of passages did not exceed 50. The cells were cultured using an established protocol, 160 with slight differences between the three laboratories, as detailed in Supplementary Data S1.

161 2.4. Standard operating protocol (SOP) for the in vitro transcriptional activation assay 162 using the UALH-hAR cell line

The following generic SOP was provided to participants laboratories as recommendations to run the assays. Due to specificities in participant laboratory facilities and uses, some slight differences between labs were noted regarding cell culture, cell exposure and luciferase readout protocols, which are detailed in Supplementary Data S1. Overall, cells were seeded in 96-wells white opaque culture plates at a density of 50,000 cells per well in culture medium. After 24 h,

cells were exposed in test medium with chemicals in a range of concentrations done in serial 168 dilutions with a factor 3.16 in DMSO (laboratories A and C) or in DMSO-containing culture 169 medium (laboratory B). In any case, the final DMSO content was always set at 0.1% v/v 170 (agonist mode) and 0.2% v/v (antagonist mode). In the antagonist mode, tested compounds 171 were exposed in presence of DHT at a concentration yielding 50-80% of the maximum 172 luciferase activity, *i.e.*, DHT 3.16 x 10⁻¹⁰ M. Each concentration was tested in technical 173 174 quadruplicates. Each plate contained the following controls: culture medium only, solvent control DMSO 0.1% v/v for agonist assays and 0.2% v/v for antagonist assays as negative 175 controls; positive control DHT 10⁻⁷ M leading to the maximal luciferase activity; in antagonist 176 assays only, the reference compound DHT 3.16 x 10⁻¹⁰ M alone yielding 50-80% of the 177 maximum luciferase activity. Plates were then incubated at 5% $CO_2 \pm 37^{\circ}C$ for 24 h. At the end 178 of the incubation, test medium was replaced with 50 µL per well of medium containing 0.5 mM 179 D-luciferin (laboratory A) or 0.3 mM D-luciferin (laboratories B and C). Luminescence signal 180 is measured in living cells for 2s per well using a luminometer. Each compound was tested in 181 at least three independent runs. 182

183 **2.5.**

Cytotoxicity assay

The viability of the cells was checked with a MTT (3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyl 184 tetrazolium bromide, Sigma M2128) assay to ensure that there was no cytotoxic effect at the 185 186 concentrations tested. After exposure, cells were incubated in 100 µL test medium containing 0.5 mg/mL MTT for 3h at 5% CO₂ and \pm 37°C. Wells were then emptied, the cells lysed in 100 187 µL DMSO/well and plates shaken for 10 minutes. The optical density (OD) was measured by 188 189 spectroscopy at 570 nm and 640 nm and OD₆₄₀ values subtracted from OD₅₇₀. Values obtained for the solvent control were considered as 100% viability. Concentrations leading to less than 190 80% viability were excluded from the analysis. 191

192 **2.6.** *Data analysis*

To assess the risk of overlap between negative and positive controls, we calculated the Z-factor, considering the basal activity in DMSO-treated cells as a negative control and the maximal activity in DHT 10⁻⁷ M-treated cells as a positive control, as follows:

$$196 Z - factor = \frac{3(standard \ deviation_{DHT \ 10-7M} + standard \ deviation_{DMSO \ control})}{|mean_{DHT \ 10-7M} - mean_{DMSO \ control}|}$$

To express luciferase activity as % of maximal effect exerted by the positive control, relative
light units (RLUs) were normalized between 0% (baseline luciferase activity in DMSO control)
and 100% (luciferase activity in DHT 10⁻⁷ M positive control) by using the following equation:

200
$$\% effect = \frac{RLU_{test\ chemical} - RLU_{DMSO\ control}}{RLU_{DHT\ 10-7M} - RLU_{DMSO\ control}}$$

A chemical is considered to have an agonistic effect when it increases luciferase activity above 201 10% of the maximum effect induced by positive control for at least two consecutive 202 203 concentrations. If not, the chemical is considered as non-agonist. A chemical is considered to have an antagonistic effect when it decreases luciferase activity by more than 30% in antagonist 204 mode. If not, the chemical is considered as non-antagonist. For active chemicals, individual 205 206 (ant-)agonist concentration-response curves are fitted with the Hill equation model, using the 207 dose-response function of a graphics and statistics software program (GraphPad Prism 6, GraphPad Software Inc.). Effective concentrations (EC) and inhibitory concentrations (IC) as 208 well the slope factor (Hill coefficient) were derived from the Hill equation. For a given 209 chemical, EC₅₀ is defined as the concentration inducing 50% of its maximal effect and IC₅₀ 210 211 represents the concentration required for 50% inhibition of the effect induced by DHT 3.16×10⁻ ¹⁰ M. Statistical analysis are performed on logEC₅₀ and logIC₅₀ values using Kruskal-Wallis 212 213 test and Dunn's multiple comparison test (p < 0.05) for the analysis of data from phases I and II, and t-test (Mann-Whitney) (p < 0.05) for the comparison of phase I with phase II. 214 To assess the WLR of the method, the CV between the three runs of a given laboratory was 215

determined by using the formula: $CV = (Standard Deviation/Mean logEC_{50} or logIC_{50}) \times 100$.

To assess the BLR of the method, we calculated the CV between the three laboratories using the formula: $CV = (Standard Deviation/Mean logEC_{50} or logIC_{50}) \times 100.$

219

3. Results and discussion

220

3.1. Phase I: Transferability of the UALH-hAR method

The phase I aimed at assessing the correct transfer of the cell line by comparing the quantification of the AR bioactivity of three reference chemicals, i.e., DHT, hydroxyflutamide and dexamethasone, chosen as reference AR agonist, AR antagonists and GR agonist, respectively.

For all three laboratories, the data produced were considered as valid. Z-factors were calculated in each laboratory and ranged from 0.7 to 0.9, meaning that the risk of overlap between negative and positive controls was negligible, as a good Z-factor should be comprised between 0.5 and 1. For all runs, the induction factor of the positive control was always more than 5 times the solvent control (from 5.6- to 31-times over all data), and the mortality never exceeded 20% (data not shown). Based on this observation, we defined a maximal basal luciferase activity of 20% and a minimal cell viability of 80% as validity criteria for the phase 2.

For all three laboratories, DHT increased luciferase activity in a concentration-dependent manner in the three independent runs as expected, with a mean logEC₅₀ of -9.8 (EC₅₀ of 1.6 x 10^{-10} M) (Fig. 2). Intra-laboratory CVs of logEC₅₀ ranged from 0.4 to 0.5% (Table 2), indicating good WLR. Regarding interlaboratory comparison, logEC₅₀ for DHT were in the same order of magnitude with a CV of 1.7%, suggesting good BLR, although there was a significant difference between laboratory A and laboratory C logEC₅₀ (Fig. 3).

In the three independent runs in all participating laboratories, HF did not induce any luciferase activity in agonist runs (data not shown) but decreased it in antagonist runs in a concentrationdependent manner, as expected (Fig. 2). The mean $\log IC_{50}$ for HF was -7.2 (IC₅₀ of 6.3 x 10⁻⁸ M). CVs in each laboratory ranged from 0.3% to 2.4%, suggesting good WLR (Table 2).
Interlaboratory CV was 2.0%, suggesting good BLR.

DEX did not display any AR-agonistic activity in any laboratory (data not shown). This lack of 243 agonistic effect of DEX in UALH-hAR cells confirmed the non-detectable expression of 244 functional GR and/or the inability of GR to induce luciferase expression from the 245 6xARE(RAD9)Luc plasmids, unlike other established models (Delfosse et al., 2012; Térouanne 246 et al., 2000; Wilson et al., 2002). However, in antagonist mode, DEX inhibited luciferase 247 activity in a concentration-dependent manner in the three laboratories (Fig. 2). The mean 248 $\log IC_{50}$ was -5.6 (IC₅₀ of 2.5 x 10⁻⁶ M) and CVs ranged from 1.0% to 3.4%, suggesting 249 250 acceptable WLR (Table 2). The mean logIC_{50s} for each laboratory were very close, with a 1.4% 251 CV, suggesting good BLR. The DEX antagonist activity observed in our study was already reported in literature in AR-CALUX, AR-EcoScreen and the mutant AR-EcoScreen GR-KO 252 cell lines (Araki et al., 2005; Sonneveld et al., 2005; Zwart et al., 2017). The fact that the UALH-253 hAR line expresses GR at very low level suggests that this effect is not due to transcriptional 254 interference with GR but rather to a direct antagonistic action of DEX on AR. Moreover, the 255 IC₅₀ of DEX on UALH-hAR was relatively high - in the µM range - while its ability to bind 256 257 to and activate human GR is reported in the nanomolar range (Molina-Molina et al., 2006). In 258 addition, binding experiments on isolated receptors have shown that DEX can bind AR at high concentrations (IC₅₀ of 1.9×10^{-4} M) (Féau et al., 2009). Further binding experiments could be 259 conducted on UALH-hAR cells using radiolabelled R1881 to assess the binding of DEX to AR 260 261 in these cells.

For all three substances, the CVs of the other Hill parameters also reflect good WLR and BLR(Supplementary Data S2).

Altogether, the three laboratories obtained perfectly concordant results from a qualitative point of view for the three substances tested. From a quantitative point of view, the logEC₅₀ and

logIC₅₀ values in the independent runs are close within and between each laboratory which indicates good WLR and BLR (Fig. 3). It is worth noting that the slight differences in the protocols between the three laboratories did not affect the comparability of the data. Main differences were related to the culture medium used (DMEM vs DMEM-F-12) and the preparation of serial dilution of chemicals (DMSO or culture medium) (Supplementary Data S1). These results show the good transferability of the method as well as its robustness when testing the three reference chemicals.

273 **3.2.** *Phase II: Assessing (anti)androgenic activity of ten substances in a blind study*

The robustness of the method was then assessed in a blind study using ten chemicals presenting different activity profiles. From a qualitative point of view, the activity profiles were perfectly consistent between the three laboratories and were all as expected for the ten substances tested (Fig. 4).

For DHT, HF and DEX, comparing logEC_{50s} and logIC_{50s} from phase I and phase II showed no statistical difference, which highlights a good robustness of the assay in quantifying the activity of these chemicals in a blinded manner within each participating laboratory.

Overall, logEC₅₀ and logIC₅₀ for active chemical substances were in the same range in the three laboratories, despite some minor but statistically significant differences between laboratories in the logEC₅₀ or logIC₅₀ of some chemicals (Fig. 5). Nevertheless, our results indicated good WLR with CVs ranging from 0.1% to 3.4%, and good BLR with CVs ranging from 1.0% to 4.0% (Table 2). The CVs of the other Hill parameters also reflect good WLR and BLR (Supplementary Data S3).

In both agonist and antagonist runs, the phase II of the study revealed good WLR and BLR,thus indicating a good robustness of the assay.

289 **3.3.** *Performance of the assay*

The activity profiles for the ten chemicals on UALH-hAR cell line were overall very 290 comparable with the ones described in the OECD Guideline 458 that uses AR-EcoScreen, AR-291 CALUX and 22Rv1/MMTV GR-KO (OECD, 2020), as well as with other similar cell models 292 (Table 3). This highlights the very good sensitivity of our method (*i.e.*, the proportion of active 293 substances that are correctly classified by the test), as activity profiles obtained for DHT, 294 R1881, testosterone, MPA, HF, BPA, M2V and BPC in UALH-hAR cells proved similar to 295 those described in the literature. For DEX, the lack of GR interference in our assay revealed a 296 direct antagonistic activity, as also reported in AR-CALUX and AR-EcoSreen-GR-KO assays. 297 298 Regarding MEHP, the lack of AR activity was previously described in similar in vitro models 299 (Stroheker et al., 2005; Takeuchi et al., 2005), although this compound alone is insufficient to conclude about specificity (the proportion of all inactive substances that are correctly classified 300 301 by the test). The antagonist effect of vinclozolin M2 is consistent with previous report in PALM cells (Molina-Molina et al., 2006) and MDA-kb2 cells (Wilson et al., 2002). In both studies, a 302 weak agonist activity of this chemical was reported. Such a weak agonistic activity was also 303 observed in our study, with a slight increase of luciferase activity by 5% at 10^{-5} M (data not 304 305 shown). However, we considered this chemical to be a full antagonist as this result does not 306 meet the criteria for agonism, *i.e.*, increase of luciferase activity above 10% for at least two 307 consecutive concentrations.

Beyond the qualitative aspect of our results, the EC₅₀ and IC₅₀ values derived from the modelled concentration-responses curves were within the ranges of acceptable values for OECD Guideline 458 for reference compounds and similar to the values found in literature (Table 3). Regarding the WLR and BLR, the CVs of logEC₅₀ and logIC₅₀ in our study were comparable to other published interlaboratory studies (Milcamps et al., 2021; Park et al., 2021).

Overall, our study suggests that the method we developed for the evaluation of AR (ant)agonistic activity of chemicals is well suited to the OECD criteria. It would be interesting to extend the screening to a higher number of chemical substances, notably regarding non active ones, to better assess the accuracy and specificity of the UALH-hAR cell line in a regulatory context.

Furthermore, a main advantage of the current assay relies on the possibility to assess 318 nonspecific effect on luciferase by using the parental UALH cell line (i.e. which expresses 319 luciferase but no functional AR) as a control for nonspecific effect on luciferase activity (Dellal 320 et al. 2020). Such a control assay is most often missing in in vitro screens based on luminescent 321 322 cell lines. Notably, it is of high relevance in the antagonist assay to distinguish between true 323 AR antagonists and chemicals that negatively interfere with luciferase activity in an ARindependent manner, e.g. through direct interference with luciferase enzyme or due to early 324 cytotoxic events. The occurrence of such nonspecific modulation of luciferase activity can also 325 be critical when assessing environmental complex mixtures such as environmental waters, 326 which anti-androgenicity is frequently detected but often at relatively high sample 327 concentrations, close to cytotoxic ones. The use of such cell control without receptor has proven 328 329 useful as quality control in environmental bioanalysis, as previously demonstrated by using the 330 UALH/UALH-hAR (Neale et al. 2020) or HG5LN/HG5LN-hPXR (Creusot et al. 2010) cell lines. 331

4. Conclusion

This interlaboratory study demonstrated the validity of the UALH-hAR-based method for the specific detection of chemicals exerting either androgenic or antiandrogenic activity. The first phase of the study demonstrated the transferability of the cell line and of the standard operating protocol to different laboratories, which appear robust as some adaptations to specific lab conditions (*e.g.*, culture medium, preparation of test chemicals) did not influence the outcome

338 of the assay, both qualitatively and quantitatively. In addition, the tests carried out under blind test conditions during the second phase produced reliable and accurate results both qualitatively 339 and quantitatively with satisfying within- and between-laboratory reproducibility of the data, 340 showing the robustness of the assay as a screening method. Comparison of the results with the 341 OECD Guideline 458 and literature data showed the performance of the assay and its relevance 342 and reliability as compared to validated AR-cell lines. Altogether, the UALH-hAR-based assay 343 represents a promising test method for the hazard assessment of chemicals potentially acting on 344 AR-signalling pathway and can be further considered for regulatory validation and international 345 346 acceptance.

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Figure 1. The three plasmid constructs transfected into U2OS to create the UALH-hAR cell
line. SV40: simian virus 40; prom: promoter; hAR: human androgen receptor; PGK:
phosphoglycerate kinase; luc: luciferase; ARE: androgen response elements from the RAD9
gene promoter; puro(R): puromycin resistance gene; hygro(R): hygromycin resistance gene;

neo(R): neomycin resistance gene; CMV: human cytomegalovirus.





Log₁₀[EC₅₀ or IC₅₀ (M)]

506 Figure 2. Phase I transactivation of human nuclear androgen receptor (hAR) by reference

ligands in UALH-hAR cells in the three laboratories. Data are means of n = 3 independent

508 experiments per laboratory; error bars represent standard deviation. Results are expressed as

the percentage of the maximum luciferase activity induced by 10^{-7} M DHT. Antagonist assays

510 are done in coexposure with 3.16 x 10^{-10} M DHT. Between experiment intra-laboratory

511 variability is illustrated in supplementary section (Figure S1).



Figure 3. Phase I comparison of logEC₅₀ and logIC₅₀ between laboratories for the testing
in triplicates of dihydrotestosterone (DHT), hydroxyflutamide (HF) and dexamethasone (DEX).
Bars represent means of 3 independent determinations. Lab: laboratory; *: p < 0.05 using

517 Kruskal-Wallis test and Dunn's multiple comparison test.

512



Figure 4. Phase II transactivation of human nuclear androgen receptor (hAR) by agonistic and antagonistic ligands in UALH-hAR cells in the three laboratories. Data are means of n =3 independent experiments per laboratory; error bars represent standard deviation. Results are expressed as the percentage of the maximum luciferase activity induced by 10⁻⁷ M DHT. Between experiment intra-laboratory variability is illustrated in supplementary section (Figures S2 and S3).



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Figure 5. Phase II comparison of logEC₅₀ and logIC₅₀ between laboratories for the blind testing of dihydrotestosterone (DHT), hydroxyflutamide (HF), dexamethasone (DEX), methyltrienolone (R1881), testosterone (T), medroxyprogesterone acetate (MPA), vinclozolin M2 (M2V), bisphenol A (BPA), bisphenol C (BPC). Bars represent means of three independent determinations. Lab: laboratory; *: p < 0.05 using Kruskal-Wallis test and Dunn's multiple comparison test.

Test chemical	Abbreviation	CAS	Chemical structure	Molecular weight (g/mol)	Purity
Dihydrotestosterone	DHT	521-18-6	H ₃ C OH H ₃ C H H ₃ C H H ₃ C OH	290.44	≥97.5%
Hydroxyflutamide	HF	52806-53-8	$H_{3}C \xrightarrow{O}_{H_{3}C OH} H \xrightarrow{O}_{CF_{3}} CF_{3}$	292.21	≥98%
Dexamethasone	DEX	50-02-2		392.46	≥98%
Methyltrienolone	R1881	965-93-5	H ₃ C CH ₃ H H H H	284.39	≥98%
Testosterone	Т	58-22-0	H ₃ C OH H ₃ C H H ₃ C H H H H	288.42	≥98%
Medroxyprogesterone acetate	MPA	71-58-9	H ₃ C H ₃ C CH ₃ H ₃ C H ₄ C CH ₃ CH ₃	386.52	≥97%
Bisphenol A	BPA	80-05-7	HO CH3	228.29	≥99%
Bisphenol C	BPC	14868-03-2	CI CI HO OH	281.13	≥98%
Vinclozolin M2	M2V	83792-61-4	H ₂ C H ₃ C OH H	260.12	≥98%
Mono(2- ethylhexyl)phthalate	MEHP	4376-20-9	OH OH CH ₃	277.34	≥97%

Table 1. Test chemicals evaluated in the interlaboratory study.

536Table 2. Activity profiles of the 10 tested chemical substances in the three laboratories in

- 537 Phase I and Phase II, represented as mean logEC₅₀ (agonist assays) and mean logIC₅₀
- 538 (antagonist assays).

			Age	onist assa	ys (logEC	50)	Anta	gonist as	says (log	IC50)
			Lab. A	Lab. B	Lab C.	Interlab.	Lab. A	Lab. B	Lab C.	Interlab.
		mean	-9.6	-9.8	-10.0	-9.8				
	Dihydrotestosterone	sd	0.04	0.05	0.05	0.17	ne	ne	ne	
		CV	0.4%	0.5%	0.5%	1.7%				
SE		mean					-7.2	-7.3	-7.1	-7.2
HAS	Hydroxyflutamide	sd	ne	ne	ne		0.02	0.18	0.15	0.15
٩		CV					0.3%	2.4%	2.2%	2.0%
		mean					-5.6	-5.7	-5.6	-5.6
	Dexamethasone	sd	ne	ne	ne		0.19	0.06	0.06	0.08
		CV					3.4%	1.0%	1.1%	1.4%
		mean	-9.8	-9.4	-9.9	-9.7				
	Dihydrotestosterone	sd	0.11	0.14	0.05	0.25	ne	ne	ne	
		CV	1.2%	1.5%	0.5%	2.6%				
		mean					-7.3	-7.2	-7.1	-7.2
	Hydroxyflutamide	sd	ne	ne	ne		0.12	0.06	0.14	0.10
		CV					1.6%	0.9%	2.0%	1.4%
	Dexamethasone	mean					-5.7	-5.8	-5.4	-5.6
		sd	ne	ne	ne		0.19	0.19	0.03	0.19
		CV	0.0	0.7	10.0	0.0	3.4%	3.3%	0.5%	3.4%
		mean	-9.8	-9.7	-10.0	- 9.8	no	no	no	
	wietnyitrienoione	su CV	0.07	0.05	0.13	0.13	ne	ne	ne	
		mean	_9.2	-9.0	-9 /	2	,			
	Testosterone	sd	0.09	0.05	0.03	- J.2 0.18	ne	ne	ne	
Щ		CV	1.0%	0.6%	0.3%	1.9%	ne	ne	ne	
IAS		mean	-8.6	-8.2	-8.8	-8.5				
Ч	Medroxyprogesterone	sd	0.08	0.02	0.09	0.34	ne	ne	ne	
	acetate	CV	0.9%	0.2%	1.0%	4.0%				
		mean								
	MEHP	sd	ne	ne	ne		ne	ne	ne	
		CV								
		mean					-7.0	-6.9	-6.7	-6.9
	Vinclozolin M2	sd	ne	ne	ne		0.13	0.07	0.08	0.17
		CV					1.8%	1.0%	1.2%	2.5%
		mean					-5.6	-5.6	-5.5	-5.6
	Bisphenol A	sd	ne	ne	ne		0.08	0.11	0.07	0.06
		CV					1.3%	2.1%	1.3%	1.0%
		mean					-6.9	-6.8	-6.5	-6.7
	Bisphenol C	sd	ne	ne	ne		0.03	0.16	0.00	0.18
		CV					0.4%	2.3%	0.1%	2.7%

Lab.: laboratory; Interlab.: interlaboratory comparison; ne: no effect; sd: standard deviation;

540 CV: coefficient of variation.

541 Table 3. Comparison of log effective concentrations logEC₅₀ and log inhibitory concentrations logIC₅₀ (log[M]) in UALH-hAR cells with

values reported in other *in vitro* reporter cell models. Bibliographic references are indicated as superscripts. *AR-EcoScreen GR-KO;
 **22Rv1/MMTV

		UALH- hAR (present study)	AR-EcoScreen	AR-CALUX	22Rv1/MMTV_GR-KO	PALM	MDA-kb2	HELN- AR(ERa-DBD)	MDA- MB453- pMMTVneo- Luc	CHO-K1 pZeoSV2AR pIND-ARE- pCMVβ-Gal
	Dihydrotestosterone	-9.7	-11.03 ~ -9.00 ^(OECD, 2020)	-9.98 ~ -9.42 ^(OECD, 2020)	-9.73 ~ -8.95 (OECD, 2020)	_9.5 (Térouanne et al., 2000)	Agonist ^(Wilson et al., 2002) -9.8 ^(Aït-Aïssa et al., 2010)			
Agonists	Methyltrienolone	-9.8	-9.6 (Araki et al., 2005)	_9.9 (Sonneveld et al., 2005)	-9.97 (Sun et al., 2016)**	-10.3 (Térouanne et al., 2000) -10.0 (Molina-Molina et al., 2006)				
	Testosterone	-9.2	-9.46 ~ -8.96 ^(OECD, 2020)	-9.25 ~ -8.8 ^(OECD, 2020)	-9.67 ~ -8.66 ^(OECD, 2020)	-7.7 (Térouanne et al., 2000)	-9.4 ^(Aït-Aïssa et al., 2010)			
	Medroxyprogesterone acetate	-8.5	-8.77 ~ -8.37 ^(OECD, 2020)	-9.23 ~ -7.75 ^(OECD, 2020)	-7.64 ~ -6.01 ^(OECD, 2020)		Agonist ^(Wilson et al., 2002)			
No effect	МЕНР	no effect							No anti-AR effect ^{(Stroheker} et al., 2005)	No (anti-)AR effect ^{(Takeuch} et al., 2005)
	Hydroxyflutamide	-7.2	-7.80 ~ -6.17 ^(OECD, 2020)	-7.80 ~ -7.54 ^(OECD, 2020)	-7.79~-7.11 ^(OECD, 2020)	Antagonist ^{(Térouanne et al.,} 2000)	Antagonist ^{(Wilson et al.,} 2002)			
	Dexamethasone	-5.6	Agonist -8.8 ^(Zwart et al., 2017) Agonist -8.6 ^(Araki et al., 2005) Antagonist -5.6 ^{(Zwart et al., 2017)*} Antagonist -4.4 ^(Araki et al., 2005)	Antagonist ^(Sonneveld et al., 2005)		Agonist ^(Térouanne et al., 2000)	Agonist ^(Wilson et al., 2002) Agonist -8.4 ^{(Aît-Aîssa et al.,} 2010)			
Antagonists	Vinclozolin M2	-6.9				Agonist -4.1 ^{(Molina-Molina et} al., 2006) Antagonist -6.8 ^{(Molina-} Molina et al., 2006)	Agonist/antagonist (Wilson et al., 2002)			
	Bisphenol A	-5.6	-7.05 ~ -4.29 ^(OECD, 2020)	-5.93 ~ -5.81 (OECD, 2020)	-5.68 ~ -5.29 ^(OECD, 2020)	Agonist -4.3 ^{(Molina-Molina} et al., 2013) Antagonist -6.0 ^{(Molina-} Molina et al., 2013)		-5.6 ^{(Delfosse et} al., 2012)		
	Bisphenol C	-6.7						-6.7 ^{(Delfosse et} al., 2012)		

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546	Supplementary Data
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548	Interlaboratory prevalidation of a new in vitro transcriptional activation assay for the
549	screening of (anti-)androgenic chemicals using the UALH-hAR cell line
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Keywords: androgen receptor; assay; stable luciferase reporter cell line; interlaboratory study.

570 Supplementary Table S1. Protocol specificities in the three laboratories.

		Laboratory A	Laboratory B	Laboratory C
Cul	ture medium	- DMEM with phenol red (Gibco 21885-025) - 5% FBS (Sigma F7524) - 1% v/v penicillin/streptomycin (Gibco 15070-63 5000 U/mL)	- DMEM/F-12 with phenol red (Gibco 31331-028) - 5% FBS (Eurobio CVFSVF00) - 1% v/v penicillin/streptomycin (Gibco 15070-63 5000 U/mL)	- DMEM/F-12 with phenol red (Gibco 31331-028) - 5% FBS (Eurobio CVFSVF00) - 1% v/v penicillin/streptomycin (Gibco 15070-63 5000 U/mL)
cu		 - 1% v/v nonessential amino acids (Gibco 11140050) - Puromycin 0.25-0.5 μg/mL (Sigma P8833) - G418 0.5-1 mg/mL (Invivogen ant-gn 100 mg/mL) - Hygromycin 0.125-0.25 mg/mL 	- Puromycin 0.25-0.5 μg/mL (Sigma P8833) - G418 0.5-1 mg/mL (Invivogen ant-gn 100 mg/mL) - Hygromycin 0.125-0.25 mg/mL	- Puromycin 0.25-0.5 μg/mL (Sigma P8833) - G418 0.5-1 mg/mL (Invivogen ant-gn 100 mg/mL) - Hygromycin 0.125-0.25 mg/mL
т	est medium	 DMEM without phenol red (Gibco 11880) 2% DCC-treated FBS 1% v/v penicillin/streptomycin (Gibco 15070-63 5000 U/mL) 1% v/v L-Glutamine 1% v/v nonessential amino acids (Gibco 11140050) 	- DMEM/F-12 without phenol red (Gibco 21041-025) - 2% DCC-treated FBS - 1% v/v penicillin/streptomycin (Gibco 15070-63 5000 U/mL) - 1% v/v L-Glutamine	 DMEM/F-12 without phenol red (Dutscher P04-41650) 2% DCC-treated FBS 1% v/v penicillin/streptomycin (Gibco 15070-63 5000 U/mL) 1% v/v L-Glutamine
Lu	iminescence medium	- Test medium with 0.5 mM D-luciferin Firefly in Tris buffer pH 8.0 (Caliper)	- Test medium with 0.3 mM D-luciferin Firefly in Tris buffer pH 8.0 (Perkin Elmer 122799)	- Test medium with 0.3 mM D-luciferin Firefly in Tris buffer pH 8.0 (Roth)
Pla	tes reference	Greiner bio-one 655098	Greiner bio-one 655083-905	Greiner bio-one 655098
Plates reference Luminometer		Synergy H4	Wallax MicroBeta TriLux	Victor3 Perkin Elmer
	Cell seeding	Cells are seeded in 96-wells white opaque culture plates at a density of 50,000 cells per well in 100 μL test medium.	Cells are seeded in 96-wells white opaque culture plates at a density of 50,000 cells per well in 150 μ L test medium.	Cells are seeded in 96-wells white opaque culture plates at a density of 50,000 cells per well in 100 μL test medium.
ı assay	Chemicals preparation	Serial dilutions of stock solutions are prepared at 1000X concentrations in DMSO and then diluted in test medium at 3X concentration.	1000X stock solution in DMSO is diluted at 4X in test medium; then, a range of 4X concentrations is done by serial dilutions in DMSO-containing test medium	Serial dilutions of stock solutions are prepared at 1000X concentrations in DMSO and then diluted in test medium at 3X concentration.
Transactivation	Exposure	50 μ L of the 3X diluted compounds are distributed to the wells containing 100 μ L test medium for a final concentration of 1X, and for a final DMSO content set at 0.1% v/v for agonist assays and 0.2% v/v for antagonist assays.	Culture medium in the plates is replaced by 150 μ L test medium then 50 μ L of the 4X concentrated compounds are added, for a final concentration of 1X, and for a final DMSO content set at 0.1% v/v for agonist assays and 0.2% v/v for antagonist assays.	50 μ L of the 3X diluted compounds are distributed to the wells containing 100 μ L test medium for a final concentration of 1X, and for a final DMSO content set at 0.1% v/v for agonist assays and 0.2% v/v for antagonist assays.
	Automation	Preparation of chemical dilutions in DMSO and their transfer into the microplates containing the cells are performed using a liquid handling workstation (Evo75, TECAN)	Cell-seeding was automated (Multidrop, ThermoFisher), and preparation of chemical dilutions and their transfer into the microplates containing the cells are performed using a liquid handling workstation (Biomek 3000, Beckman).	All the steps are performed manually.

			Dihydrotest	osterone		H	lydroxyflutamic	le	Dexamethasone		
		Hill slope	logEC ₁₀	logEC ₅₀	Max (%)	Hill slope	logIC ₃₀	logIC ₅₀	Hill slope	logIC ₃₀	logIC ₅₀
	Run 1	1.7	-10.2	-9.6	99.1	-0.9	-7.6	-7.2	-0.9	-6.2	-5.8
	Run 2	2.0	-10.1	-9.6	97.2	-1.1	-7.6	-7.2	-1.0	-5.7	-5.4
Laboratory A	Run 3	2.5	-10.0	-9.7	97.7	-1.3	-7.5	-7.2	-0.9	-6.0	-5.6
Laboratory A	mean	2.1	-10.1	-9.6	98.0	-1.1	-7.6	-7.2	-0.9	-6.0	-5.6
	sd	0.37	0.08	0.04	1.03	0.17	0.08	0.02	0.05	0.21	0.19
	CV	17.9%	0.8%	0.4%	1.1%	15.7%	1.0%	0.3%	5.7%	3.6%	3.4%
	Run 1	1.7	-10.4	-9.8	101.4	-1.1	-7.5	-7.2	-0.8	-6.1	-5.7
	Run 2	1.4	-10.5	-9.8	99.6	-0.8	-8.0	-7.5	-0.9	-6.1	-5.7
Laboratory P	Run 3	1.4	-10.4	-9.7	98.9	-1.0	-7.7	-7.3	-0.8	-6.2	-5.8
Laboratory B	mean	1.5	-10.4	-9.8	99.9	-1.0	-7.7	-7.3	-0.8	-6.1	-5.7
	sd	0.19	0.06	0.05	1.31	0.14	0.23	0.18	0.04	0.08	0.06
	CV	12.8%	0.6%	0.5%	1.3%	14.1%	3.0%	2.4%	4.9%	1.2%	1.0%
	Run 1	1.7	-10.4	-10.0	98.8	-0.9	-7.3	-6.9	-0.8	-6.0	-5.5
	Run 2	1.9	-10.5	-9.9	103.4	-1.1	-7.4	-7.1	-0.8	-6.0	-5.6
Laboratory C	Run 3	2.4	-10.4	-10.0	99.5	-0.8	-7.7	-7.2	-0.8	-6.0	-5.6
Laboratory C	mean	2.0	-10.4	-10.0	100.6	-0.9	-7.5	-7.1	-0.8	-6.0	-5.6
	sd	0.35	0.06	0.05	2.47	0.15	0.18	0.15	0.04	0.04	0.06
	CV	17.6%	0.6%	0.5%	2.5%	16.3%	2.5%	2.2%	4.8%	0.6%	1.1%
Interleherater	mean	1.9	-10.3	-9.8	99.5	-1.0	-7.6	-7.2	-0.9	-6.0	-5.6
mean	sd	0.30	0.19	0.17	1.36	0.11	0.13	0.15	0.08	0.09	0.08
	CV	16.2%	1.8%	1.7%	1.4%	11.1%	1.7%	2.0%	9.0%	1.5%	1.4%

572 Supplementary Table S2. Phase I intra- and interlaboratory comparisons of the effects of dihydrotestosterone, hydroxyflutamide and

573	dexamethasone based	on the Hill model	parameters: H	Hill slope, I	$logEC_{10}$,	logEC50 and	maximum o	or logIC ₃₀ and	logIC ₅₀ .
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574 Sd: standard deviation; CV: coefficient of variation; Max: maximum luminescence reached by the tested compound (% of max induced by DHT 10^{-7} M).

		Dihydrotestosterone				н	ydroxyflutamid	e	Dexamethasone			
		Hill slope	logEC ₁₀	logEC ₅₀	Max (%)	Hill slope	logIC ₃₀	logIC₅0	Hill slope	logIC ₃₀	logIC ₅₀	
	Run 1	1.6	-10.3	-9.7	111.5	-1.3	-7.5	-7.2	-1.6	-5.8	-5.6	
	Run 2	2.1	-10.4	-9.9	107.4	-1.3	-7.7	-7.4	-1.3	-6.0	-5.7	
Lohoustow. A	Run 3	1.4	-10.4	-9.7	104.6	-1.1	-7.6	-7.3	-0.9	-6.4	-5.9	
Laboratory A	mean	1.7	-10.4	-9.8	107.8	-1.3	-7.6	-7.3	-1.2	-6.0	-5.7	
	sd	0.39	0.07	0.11	3.47	0.14	0.11	0.12	0.35	0.29	0.19	
	CV	22.7%	0.6%	1.2%	3.2%	10.8%	1.5%	1.6%	28.5%	4.8%	3.4%	
	Run 1	1.2	-10.3	-9.5	99.1	-1.1	-7.4	-7.1	-0.7	-6.4	-5.9	
	Run 2	1.1	-10.1	-9.3	101.1	-1.0	-7.5	-7.2	-0.7	-6.4	-5.9	
lekenstern D	Run 3	1.1	-10.4	-9.5	101.8	-1.0	-7.6	-7.2	-1.3	-5.9	-5.6	
Laboratory B	mean	1.2	-10.2	-9.4	100.7	-1.1	-7.5	-7.2	-0.9	-6.2	-5.8	
	sd	0.06	0.14	0.14	1.38	0.05	0.08	0.06	0.34	0.34	0.19	
	CV	5.0%	1.4%	1.5%	1.4%	4.7%	1.0%	0.9%	38.9%	5.4%	3.3%	
	Run 1	2.3	-10.4	-10.0	99.2	-0.9	-7.4	-6.9	-1.0	-5.8	-5.5	
	Run 2	1.7	-10.4	-9.9	103.1	-1.0	-7.5	-7.1	-1.0	-5.8	-5.4	
laboratory C	Run 3	1.7	-10.4	-9.9	118.6	-1.0	-7.6	-7.2	-0.9	-5.8	-5.4	
Laboratory C	mean	1.9	-10.4	-9.9	107.0	-0.9	-7.5	-7.1	-1.0	-5.8	-5.4	
	sd	0.36	0.04	0.05	10.25	0.05	0.12	0.14	0.04	0.03	0.03	
	CV	18.7%	0.3%	0.5%	9.6%	5.3%	1.7%	2.0%	3.7%	0.6%	0.5%	
late debe vetera.	mean	1.6	-10.3	-9.7	105.2	-1.1	-7.5	-7.2	-1.0	-6.0	-5.6	
mean	sd	0.38	0.08	0.25	3.91	0.16	0.05	0.10	0.19	0.21	0.19	
mean	CV	24.3%	0.8%	2.6%	3.7%	14.8%	0.7%	1.4%	18.8%	3.5%	3.4%	

576 Supplementary Table S3. Phase II intra- and interlaboratory comparisons of the effects of dihydrotestosterone, hydroxyflutamide and

577	dexamethasone based	on the Hill model parameter	s: Hill slope, logEC ₁₀ , logEC ₅₀ and	maximum or logIC ₃₀ and logIC ₅₀ .
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578 Sd: standard deviation; CV: coefficient of variation; Max: maximum luminescence reached by the tested compound (% of max induced by DHT 10^{-7} M).

			Methyltri	enolone			Testosterone				Medroxyprogesterone acetate			
		Hill slope	logEC ₁₀	logEC ₅₀	Max (%)	Hill slope	logEC ₁₀	logEC ₅₀	Max (%)	Hill slope	logEC ₁₀	logEC ₅₀	Max (%)	
	Run 1	1.5	-10.4	-9.8	98.6	1.5	-9.8	-9.2	87.3	1.9	-9.1	-8.6	46.6	
	Run 2	1.5	-10.3	-9.7	94.4	1.4	-10.0	-9.3	84.1	1.9	-9.1	-8.6	42.2	
Laboratory A	Run 3	1.3	-10.5	-9.8	91.4	1.6	-9.7	-9.1	91.2	1.4	-9.2	-8.5	51.8	
Laboratory A	mean	1.5	-10.4	-9.8	94.8	1.5	-9.9	-9.2	87.5	1.7	-9.2	-8.6	46.9	
	sd	0.14	0.13	0.07	3.61	0.10	0.13	0.09	3.55	0.33	0.04	0.08	4.80	
	CV	9.5%	1.2%	0.8%	3.8%	6.7%	1.4%	1.0%	4.1%	18.8%	0.5%	0.9%	10.2%	
	Run 1	1.0	-10.7	-9.8	101.3	1.2	-9.9	-9.1	95.5	1.9	-8.7	-8.2	58.9	
	Run 2	1.2	-10.4	-9.7	98.1	1.1	-9.9	-9.0	96.6	1.3	-8.9	-8.1	62.2	
Laboratory B	Run 3	1.4	-10.4	-9.7	102.3	1.0	-10.0	-9.0	97.8	1.3	-8.9	-8.2	62.1	
Laboratory B	mean	1.2	-10.5	-9.7	100.6	1.1	-9.9	-9.0	96.6	1.5	-8.8	-8.2	61.1	
	sd	0.18	0.14	0.05	2.21	0.09	0.03	0.05	1.13	0.34	0.14	0.02	1.91	
	CV	14.8%	1.4%	0.5%	2.2%	8.1%	0.3%	0.6%	1.2%	22.6%	1.6%	0.2%	3.1%	
	Run 1	2.0	-10.4	-9.9	93.9	1.4	-10.0	-9.4	104.9	1.3	-9.5	-8.7	58.9	
	Run 2	1.6	-10.7	-10.1	103.0	1.2	-10.2	-9.4	96.4	1.1	-9.8	-8.9	52.2	
Laboratory C	Run 3	1.6	-10.5	-9.9	108.1	1.1	-10.3	-9.4	103.3	1.5	-9.5	-8.8	54.7	
Laboratory C	mean	1.8	-10.5	-10.0	101.7	1.3	-10.2	-9.4	101.5	1.3	-9.6	-8.8	55.3	
	sd	0.22	0.16	0.13	7.17	0.16	0.11	0.03	4.54	0.20	0.17	0.09	3.39	
	CV	12.6%	1.5%	1.3%	7.1%	12.3%	1.0%	0.3%	4.5%	14.9%	1.8%	1.0%	6.1%	
Interlahoratory	mean	1.5	-10.5	-9.8	99.0	1.3	-10.0	-9.2	95.2	1.5	-9.2	-8.5	54.4	
mean	sd	0.27	0.06	0.13	3.68	0.21	0.15	0.18	7.10	0.20	0.37	0.34	7.14	
mean	CV	18.0%	0.6%	1.4%	3.7%	16.2%	1.5%	1.9%	7.5%	13.4%	4.1%	4.0%	13.1%	

580 Supplementary Table S4. Phase II intra- and interlaboratory comparisons of the effects of methyltrienolone, testosterone and

581 **medroxyprogesterone acetate** based on the Hill model parameters: Hill slope, logEC₁₀, logEC₅₀ and maximum.

582 Sd: standard deviation; CV: coefficient of variation; Max: maximum luminescence reached by the tested compound (% of max induced by DHT
 583 10⁻⁷ M).

Supplementary Table S5. Phase II intra- and interlaboratory comparisons of the effects of vinclozolin M2, bisphenol A and bisphenol C based on the Hill model parameters: Hill slope, logIC₃₀ and logIC₅₀.

		Vi	nclozolin M2			Bisphenol A			Bisphenol C			
		Hill slope	logIC ₃₀	logIC₅₀	Hill slope	logIC ₃₀	logIC ₅₀	Hill slope	logIC ₃₀	logIC ₅₀		
	Run 1	-1.0	-7.4	-7.0	-1.0	-5.9	-5.6	-1.8	-7.0	-6.8		
	Run 2	-1.2	-7.3	-7.0	-1.4	-6.0	-5.7	-1.3	-7.1	-6.9		
Laboratory A	Run 3	-1.0	-7.6	-7.2	-0.8	-6.1	-5.6	-1.4	-7.1	-6.9		
Laboratory A	mean	-1.1	-7.4	-7.0	-1.1	-6.0	-5.6	-1.5	-7.1	-6.9		
	sd	0.15	0.16	0.13	0.32	0.07	0.08	0.27	0.07	0.03		
	CV	13.9%	2.2%	1.8%	29.6%	1.2%	1.3%	17.5%	1.0%	0.4%		
	Run 1	-1.0	-7.3	-7.0	-1.0	-6.0	-5.6	-0.9	-7.3	-6.9		
	Run 2	-1.1	-7.1	-6.8	-0.9	-6.1	-5.6	-0.8	-7.3	-6.8		
Laboratory P	Run 3	-1.3	-7.1	-6.9	-1.0	-5.8	-5.4	-0.9	-7.0	-6.6		
Laboratory B	mean	-1.1	-7.2	-6.9	-0.9	-5.9	-5.6	-0.9	-7.2	-6.8		
	sd	0.20	0.12	0.07	0.07	0.13	0.11	0.04	0.16	0.16		
	CV	17.4%	1.6%	1.0%	7.1%	2.3%	2.1%	4.4%	2.3%	2.3%		
	Run 1	-1.0	-7.1	-6.7	-0.8	-6.1	-5.6	-1.0	-6.9	-6.5		
	Run 2	-0.9	-7.2	-6.8	-0.8	-5.9	-5.5	-1.0	-6.9	-6.5		
Laboratory C	Run 3	-1.1	-7.0	-6.6	-0.9	-5.9	-5.5	-1.5	-6.8	-6.5		
Laboratory C	mean	-1.0	-7.1	-6.7	-0.8	-6.0	-5.5	-1.2	-6.8	-6.5		
	sd	0.06	0.10	0.08	0.07	0.10	0.07	0.28	0.06	0.005		
Interlaboratory mean	CV	6.4%	1.4%	1.2%	8.5%	1.6%	1.3%	23.8%	0.9%	0.1%		
	mean	-1.1	-7.2	-6.9	-0.9	-6.0	-5.6	-1.2	-7.0	-6.7		
	sd	0.07	0.16	0.17	0.13	0.02	0.06	0.34	0.18	0.18		
	CV	6.4%	2.2%	2.5%	13.4%	0.4%	1.0%	28.5%	2.6%	2.7%		

586 Sd: standard deviation; CV: coefficient of variation.



588

Log₁₀[EC₅₀ or IC₅₀ (M)]

Figure S1. Phase I transactivation of human nuclear androgen receptor (hAR) by 589

590 **reference ligands** in UALH-hAR cells : intra-laboratory variability. n = 3 independent

experiments. Results are expressed as the percentage of the maximum luciferase activity 591

induced by 10⁻⁷ M DHT. Antagonist assays are done in coexposure with 3.16 x 10⁻¹⁰ M DHT. 592



594

Log₁₀[EC₅₀ (M)]

Figure S2. Phase II transactivation of human nuclear androgen receptor (hAR) by agonistic ligands in UALH-hAR cells: intra-laboratory variability in the three laboratories. n = 3 independent experiments per laboratory. Results are expressed as the percentage of the maximum luciferase activity induced by 10^{-7} M DHT.



Figure S3. Phase II transactivation of human nuclear androgen receptor (hAR) by antagonistic ligands in UALH-hAR cells: intra-laboratory variability in the three laboratories. n = 3 independent experiments. Results are expressed as the percentage of the maximum luciferase activity induced by 10^{-7} M DHT. Antagonist assays are done in coexposure with 3.16×10^{-10} M DHT.