

Various Genital and Reproductive Phenotypes in 46,XX/46,XY Chimeras

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Tetragametic chimeras are due to the fusion of two different zygotes after fertilization. When occurring between embryos of different chromosomal sex, the phenotype ranges from fertile individuals to infertile patients and even to patients with variations in sex development. Here we report three new cases of XX/XY chimeras, one in a young boy carrying an abnormal gonad which turned out to be an ovary and two in phenotypically normal infertile men, one of whom had been diagnosed previously as a XX-SRY negative male. These cases highlight the importance of combining several cytogenetic and molecular techniques on different tissues for a proper diagnosis and an appropriate prognosis.

Introduction:

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59 Chimera and chimerism are defined by the coexistence of two whole distinct genomes within the 60 same organism. They differ from mosaicism where the organism contains only one genome where a mutation or a chromosomal event occurred during the first mitotic divisions of the zygote, leading 61 62 to two different cellular populations. Chimerism is confined to one tissue or to one compartment. It can be temporary in the course of 63 64 blood transfusion or definitive as the consequence of allografting in patients receiving an organ or 65 bone marrow cells from a HLA-compatible donor. Chimerism can also be the result of hematopoietic stem cell travel from an embryo to another during dizygotic or even monozygotic 66 twin pregnancy (Bourthoumieu et al., 2005, 2006). This type of chimerism must be distinguished 67 68 from microchimerism, a normal process during every pregnancy which is due to the bi-directional trafficking of fetal and maternal cells through placenta. The persistence of fetal cells in pregnant 69 70 women, and even in mothers a long time after the birth of their child, predispose them to develop 71 various autoimmune diseases, particularly multiple sclerosis (Aractingi et al., 1998; Nelson, 2002; 72 Bloch et al., 2011) 73 In humans, chimeras generally occur after the fusion of two different zygotes, thus resulting in a 74 single but dizygotic embryo named tetragametic chimera (Wenk, 2018). In vitro fertilization, and the replacement of more than one embryo, has been suspected to increase this risk of fusion 75 76 (Simon-Bouy et al., 2003). In some rare cases, accidental fertilization of a second polar body (Chen 77 et al. 2005) or parthenogenetic development of a zygote can lead to the formation of a chimera 78 (Strain et al, 1995, Kaise-Roger et al. 2006, Conlin et al., 2010). 79 Whether a chimera results from the fusion of two zygotes with the same (XX/XX or XY/XY) or 80 different (XX/XY) chromosomal genders, and depending on the repartition of each genome within 81 the different tissues, the phenotype range from strictly typical and fertile male or female individuals 82 (Bromilow and Duguid, 1989) to infertile patients (Laursen et al., 2018, Choi et al., 2019) and even 83 to a complete absence of sexual differentiation (Sugawara et al., 2012). Here we report three new

cases of XX/XY chimeras in male patients who have been investigated by using conventional and molecular cytogenetic or genetic techniques. These cases raise the possibility that some XX-SRY negative males, some infertile patients or some children with a variation in sex development could be in fact chimeras.

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Patients

- Patients 1 and 2 and the parents of patient 3 gave their written informed consent and signed for
- 91 genetic analyses performed according to the rules of the Agence de la Biomédecine (France)
- 92 according to the French legislation.
- 93 Patient 1
- Patient 1 was a 26 years old male consulting for infertility. Clinical examination showed a normal
- penis with two intrascrotal testes measuring 10 mL on both sides.
- 96 Endocrine analysis showed normal inhibin B (86 pg/ml), FSH (4,7 IU/l) and testosterone 4ng/ml)
- 97 levels.
- 98 Karyotype analysis was initially performed from blood lymphocytes using standard techniques and
- 99 R and G banding. It showed an homogenous 46,XX chromosomal female constitution confirmed by
- 100 FISH using chromosome X and Y centromeric probes on 400 interphase nuclei and by FISH with a
- probe coding for the SRY gene which showed that this gene was not present on the X chromosomes.
- 102 SNP-array analysis (OmniExpress-24, 715K, Illumina) showed a normal 46, XX profile without
- any deletion or duplication, especially upstream from the SOX9 gene (Benko et al., 2011). The total
- lack of Y chromosome in peripheral blood was also confirmed by PCR analysis which showed the
- absence of SRY and ZFY sequences as well as the total lack of AZFa, b and c regions (Figure 1).
- 106 Therefore, this patient was initially diagnosed as a XX SRY-negative male. However, sperm
- analysis revealed a cryptozoospermia at the first examination and then a severe oligozoospermia
- 108 (20 000 spermatozoa/ml) which allowed the cryopreservation of sperm samples. Because such a
- 109 result is theoretically impossible in XX testicular DSD (Disorders/Differences of Sex

Development), Y-positive cells were sought in another tissue than blood and, indeed, FISH analysis with chromosome X and Y centromeric probes revealed 53% of XY cells on a buccal cell smear, mixed with XX cells (Figure 1). Unfortunately, genomic DNA extraction from buccal cells failed.

Sperm FISH analysis using chromosomes X, Y and 18 centromeric probes was then performed on a sample of ejaculated sperm cells (n=50). It showed a normal ratio 1:1 between X and Y bearing spermatozoa thus indicating that these gametes were produced from normal XY meiotic cells.

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Patient 2

Patient 2 was a 35 years old male consulting for infertility. Clinical examination showed a normal penis with two intrascrotal testes measuring 3.5 ml on the left and 8 ml on the right. The left testis also presented a stage 3 varicocele. Sperm analysis revealed a non-obstructive azoospermia and endocrine analysis showed decreased inhibin (18pg/ml) and increased FSH (16 IU/l) levels and a normal testosterone level (3.68 ng/ml). Karyotype study was performed from blood lymphocytes using standard techniques and R and G banding. It showed the presence of two different cell lines: one with a normal 46,XY male karyotype in 75% of the cells analyzed and one with a normal 46,XX female karyotype in the others. Molecular analysis of the Y chromosome using PCR was normal indicating the presence of the SRY gene as well as the lack of deletion of the AZFa, b and c regions. The two populations were also present in the same proportions (75% XY, 25% XX), on cells from a buccal swab analyzed by FISH using probes coding for the X and for the Y centromeres. A SNP-array analysis (OmniExpress-24, 715K, Illumina) was performed on DNA extracted from blood lymphocytes. The high number of SNPs on this type of array (more than 700 000) allowed identifying 9 distinct allele frequencies confirming the presence of 4 different haplotypes in a mosaic state (Figure 2). This patient underwent a testicular biopsy which revealed a normal testicular architecture associated with Leydig cell hyperplasia in the right gonad. Sperm cells were successfully retrieved and frozen. Sperm FISH analysis using chromosomes X, Y and 18 centromeric probes was performed on the

supernatant obtained after biopsy shredding. It showed a normal ratio 1:1 between X and Y bearing spermatozoa thus indicating, like in patient 1, that these gametes were produced from normal XY meiotic cells (Figure 1). FISH study of paraffin embedded sections using the same probes showed a majority of XY cells in both gonads with around 80% of Y-positive nuclei and 20% of XX cells. Most of these XX cells were found in the interstitial compartment of the testis and in the peritubular sheath of seminiferous tubules. However, some of them were also found within the seminiferous tubules (Figure 1).

The couple underwent *in vitro* fertilization after intra-cytoplasmic sperm injection which resulted in a singleton pregnancy and in the birth of a healthy boy.

Patient 3

He was a 9 years old boy referred for clinical and biological follow-up after left gonadectomy. It was performed at the age of six years in the course of the chirurgical cure of unilateral cryptorchidism because of the abnormal aspect of the gonad. Histological examination of this latter showed a gonad corresponding to an ovary with primordial follicles surrounded by a canal structure evoking Mullerian remains like a hemi-uterus. Sections of tubes corresponding either to a hypoplastic epididymis or to Wolffian vestigial structures were also described. No testicular tissue was observed. At the time of his evaluation, clinical examination showed a well-developed boy weighing 29.2kgs (+0.5 SD) and measuring 1.28m. Left testis was lacking and the right one was in scrotal position, measuring 15x10mm. Penis was 45mm long. Achromic spots were noticed on the whole body. Chromosomal analysis on blood cells after R and G banding revealed a 46,XX female karyotype in 76% of cells and a 46,XY male karyotype in 24%. These percentages were slightly different on cells from a buccal swab (60% XX, 40% XY) analyzed by FISH with chromosome X and Y centromeric probes. SNP-array analysis (Humanomniexpress-24 715K, Illumina) performed on

DNA from blood cells confirmed the diagnosis of tetragametic chimera by visualizing nine different

allele frequencies and estimating the rate of XY cells to 10-15%.

FISH study of paraffin embedded sections of the removed gonad using the same probes showed

percentages of XY ranging from 4% to 90% according to the type of histological structure (4% in

the ciliated epithelium vs 90% in the epididymis-like tubes) (Figure 3).

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Discussion:

As for the diagnosis and the evaluation of mosaicism, chromosomal analysis by FISH is a powerful and sensitive method for detecting and measuring chimera. It allows a cell by cell identification of the chromosomal content as opposed to molecular techniques which give a result as a whole. In the 3 patients we analyzed, FISH was able to give a precise evaluation of the XY/XX cell ratio and, when coupled to morphological study on histological tissue sections, to correlate this ratio to the nature of the tissue or the cells like mature gametes in patients 1 and 2. On the contrary, the very high sensitivity of PCR allowed us to assert that patient 1 had a homogenous XX chromosomal constitution in blood cells with absolutely no Y chromosome material. In patient 2, PCR indicated also that his Y chromosome had a normal structure, without any molecular deletion. Therefore, we preconize that patients suspect to carry a chromosomal mosaicism or to be a chimera should be studied by using both cytogenetic, including FISH, and molecular techniques. Such a procedure has been particularly useful in patient 1 who was initially considered as a 46,XX-SRY negative male. The fact that he presented with cryptozoospermia led us to question this diagnosis and to perform a FISH analysis on another tissue than blood. A smear of cells from a buccal swab appeared to be the simplest and non-invasive technique for exploring a chromosomal mosaicism or a chimera by FISH and, in his case, it allowed to correct the diagnosis. If azoospermic, this patient certainly would not have been evaluated by this technique. This raises the question of the diagnostic procedure in XX-SRY negative males in whom molecular investigations fail to find a cause such as a molecular duplication upstream from SOX9 gene (Benko et al, 2011) or a mutation in the WNT4/RSPO1 gene

187 pathway (Micali et al., 2005). Therefore, we recommend that every patient in this situation should 188 have a complementary examination of another tissue than blood, by FISH or PCR, in order to detect 189 a possible XX/XY chimera. 190 In our patients, FISH was also applied on testicular cells and mature sperm cells when available. In 191 patients 1 and 2, it revealed a normal 1:1 ratio between X and Y bearing spermatozoa confirming 192 that spermatogenesis is possible only from XY spermatogonia and that meiotic segregation of sex 193 chromosomes at meiosis is normal, even in the presence of a variable contingent of XX cells. To 194 our knowledge, only one study described the evaluation of testicular cells by FISH in a 195 46,XX/46,XY true hermaphrodite patient (Sugawara et al., 2012) but the authors presented their 196 results as a whole without discriminating spermatozoa from other testicular cells. Our results are of 197 importance since they suggest strongly that spermatozoa from XX/XY males are chromosomally 198 normal and can be used by in vitro fertilization to give birth to a normal offspring, as it was the case 199 in patient 2. 200 Finding XX cells within seminiferous tubules raises the question of their origin. Since the 201 differentiation of supporting cells into Sertoli cells in the primitive gonad needs the expression of 202 the Sry gene in mouse (Ungewitter and Yao, 2013), XX cells in a human adult seminiferous tubule, 203 as found in patient 2, are unlikely to be Sertoli cells. Therefore, they might be germ cells which would be unable to initiate the meiotic process although it remains to be elucidated. 204 205 In patient 3, who was still a child, FISH was only possible on paraffin-embedded sections from the 206 unilateral gonadectomy that he underwent at the age of six. However, results were particularly 207 interesting since they showed that most of the XY cells were localized in the male-differentiated 208 parts of the gonad, i.e. those resembling to epididymis-like tubes, 90% vs 4% in the ciliated 209 structures probably deriving from a uterine horn (Figure 3). This observation highlights the 210 usefulness of FISH performed on histological sections which can superimpose genetic data to tissue 211 or cellular structures.

212 Comparative genomic hybridization on DNA-arrays (CGH-array) is a very sensitive method for the 213 detection of chromosomal imbalances but is unable to diagnose balanced structural chromosomal 214 rearrangements. When performed with SNP-arrays, it allows determining the genotype of the 215 patients by analyzing the status of each DNA marker designed as the B-allele frequency (BAF): 216 homozygous (AA or BB) or heterozygous (AB), leading to the visualization of three dot plots on 217 the graph. Therefore, SNP-array can detect losses of heterozygosity of entire chromosomal regions, 218 as seen in consanguineous patients, as well as uniparental isodisomy or heterodisomy, when 219 parental DNAs are available, in patients presenting with a genetic disease due to parental genomic 220 imprinting (Conlin et al., 2010; Kearney et al, 2011). Moreover, SNP-array is also able to diagnose 221 a maternal contamination in fetal cell samples by revealing more than two different haplotypes. 222 Indeed, a 50/50 mix of two different DNAs would lead to six recognizable haplotypes (AA/AA, AA/AB or AB/AA, AA/BB or BB/AA, AB/AB, AB/BB or BB/AB and BB/BB). The fact that 223 224 DNAs are in different proportions reaches this number to nine and the frequency of each haplotype 225 gives the level of contaminating DNA. In chimeras due to the fusion of two embryos, the situation 226 is stackable to that seen in the contamination of a DNA sample by an external one. In patient 1, 227 SNP-array performed on blood cells was normal since he had an homogenous 46,XX constitution in 228 this tissue and DNA from a buccal swab was not available. In patient 2 and 3, SNP-array profile revealed nine dot plots confirming that both chimeras had a tetragametic origin. 229 230 Another explanation to the homogenous XX karyotype in patient 1 could be a chimerism secondary to blood exchange with a female fetus during pregnancy, as already described (Sudik et al., 2001; 231 232 Johannsen et al., 2003). To our knowledge, in patient 1, pregnancy was not twin. This information 233 should be systematically sought in case of discordance between blood karyotype and phenotype in 234 order to avoid confusion between chimerism and disorder/difference of sex development. 235 In conclusion, these three reports illustrate the genital phenotype of XX/XY chimera in man when 236 abnormal. They highlights situations where the search for chimera, or chimerism, should be 237 performed i) in 46,XX-SRY negative males in whom known molecular causes have been excluded

ii) in men presenting with idiopathic non-obstructive infertility iii) in children with a 238 239 difference/disorder of sex development without cytogenetic or molecular cause identified. Diagnosis of chimera should be made on different tissues by conventional and molecular (FISH) 240 241 techniques completed by a SNP-array study in order to identify the causal mechanism. 242 243 Author's roles: AH, AV, SCB, MP and HM made the study and wrote the paper. EA and SB realized the clinical 244 245 and biological care of patient 1. XF, JFH and RBB realized the clinical and biological care of patient 2. CA, ACT, IGV and FD realized the clinical and biological care of patient 3. 246 247 Statement of Ethics: Patients or parents of patients gave their written informed consent for the

genetic analyses performed in this study. The research was conducted ethically in accordance with

the Word Medical Association Declaration of Helsinki.

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251 Conflict of interest: The authors have no conflicts of interest to declare.

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317 **Legends**

- 318 <u>Figure 1</u>:
- Patient 1: A) FISH on blood cells with probes coding for the chromosome X centromere and for the
- 320 SRY gene showed a homogenous XX profile and the lack of SRY at the tip of one of the X
- 321 chromosomes. B) PCR on blood cell DNA with primers specific of the AZF regions of the Y
- 322 chromosome. Patient's DNA (lane 4) gave a negative result as for female DNA (lane 5) when
- 323 compared to male patients (lane 2 and 3). C) SNP-array on blood cell DNA revealed a normal X
- 324 chromosome ratio when compared to XX DNA (bottom). B-allele frequency (top) showed only
- 325 three dot plots indicating the existence of two different haplotypes. D) FISH on buccal cells with
- probes coding for the chromosomes X (green) and Y (red) centromeres found 53% of XY cells.
- 327 Figure 2:
- Patient 2: A and B) FISH on buccal cells (A) and blood cells (B) with probes coding for the
- 329 chromosomes X (green) and Y (red) centromeres showed 25% of XX cells and 75% of XY cells. C)
- 330 SNP-array on blood cell DNA. The B-allele frequency revealed nine different haplotypes indicating
- the existence of a tetragametic chimera. D) FISH on the supernatant of the testicular biopsy with
- probes coding for the chromosomes X (green) and Y (red) centromeres showed both XX and XY
- cells (fine arrows) mixed with haploid X or Y positive cells with a smaller nucleus (large arrows).
- E) FISH on paraffin embedded sections with probes coding for the chromosomes X (green) and Y
- 335 (red) centromeres showing the presence of XX cells in the peritubular sheath of seminiferous
- tubules. F) On the same sections, rare XX cells are also found within seminiferous tubules. The
- 337 peritubular sheath is visible on the right as a faint green fluorescence containing an elongate
- 338 nucleus.
- 339 Figure 3:
- Patient 3: FISH on paraffin embedded sections of the gonadectomy with probes coding for the
- 341 chromosomes X (green) and Y (red) centromeres. Most nuclei in the epithelium surrounding the

- 342 tubules corresponding to a hypoplastic epididymis were XY (A) while most cells in the rest of the
- gonad corresponding to an ovary and to tubules evoking Mullerian remains were XX (B).

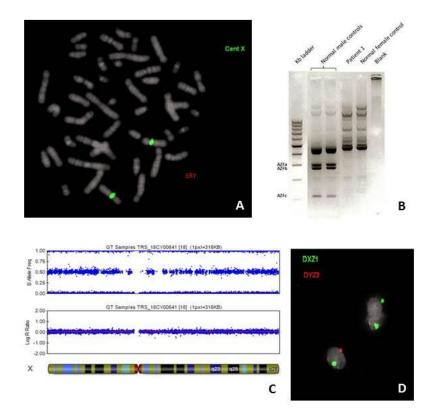


Figure 1

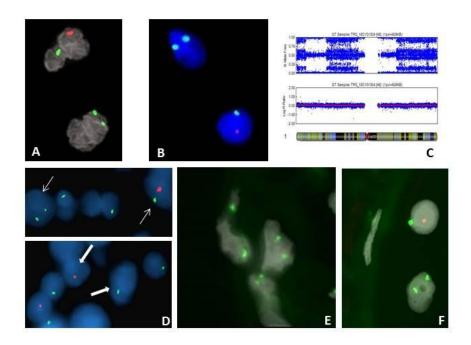
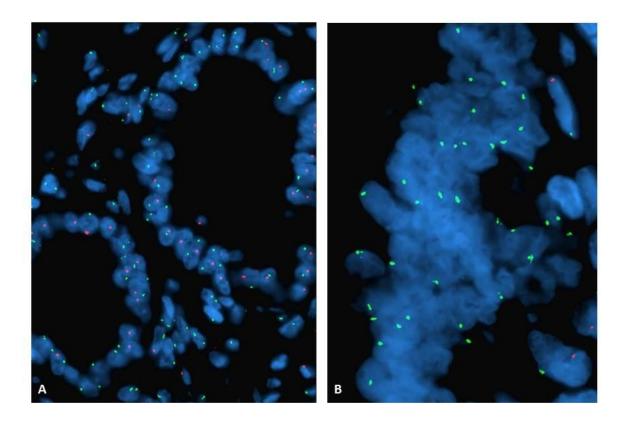


Figure 2



351 Figure 3