

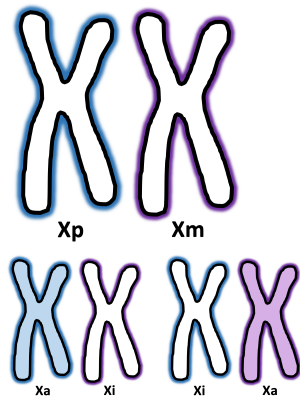
**X Chromosome Inactivation (XCI):** The epigenetic process which silences one X chromosome in females to achieve gene dosage balance between males and females. [1]

**Xa:** the active X chromosome

**Xi:** the inactive X chromosome

**Xm:** maternally inherited X chromosome

**Xp:** paternally inherited X chromosome



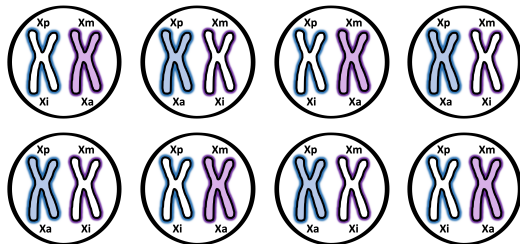
*Figure 1: The paternally inherited (Xp) chromosome in and the maternally inherited chromosome (Xm). The active chromosome Xa is represented with color filled in, while the inactive chromosome Xi is represented with no color fill.*

**Escape:** The status of genes which are expressed on Xi. These genes “escape” inactivation.

**Inactive:** The status of genes which are suppressed on Xi.

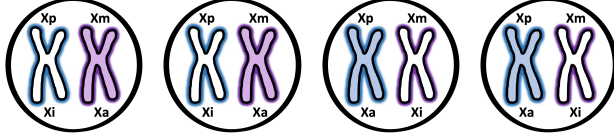
**Variable:** The status of genes which are expressed on Xi in some tissues/individuals, but not in all tissues/individuals. [2]

**Mosaicism:** Xa and Xi are typically randomly assigned to either Xp or Xm in each cell. *Mosaicism* refers to the heterogeneity of Xa/Xi assignment among a set of cells.

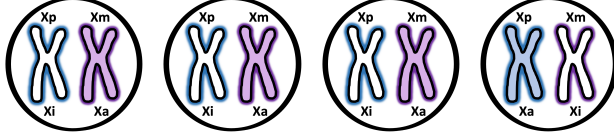


*Figure 2: Mosaicism. A sample of six cells with differences in Xa/Xi assignments.*

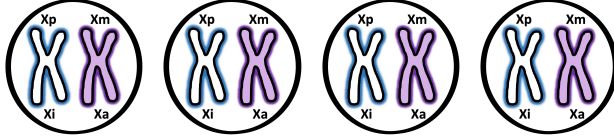
**Skew:** A ratio which represents the mosaicism of Xa/Xi assignment in a set of cells.



**Figure 3a: Skew Ratio 1.** Two samples have  $X_i$  assignment on  $X_m$ , while two samples have  $X_i$  assignment on  $X_p$ . The resulting skew ratio is 50:50.



**Figure 3b: Skew Ratio 2.** Three samples have  $X_i$  assignment on  $X_m$ , while one sample has  $X_i$  assignment on  $X_p$ . The resulting skew ratio is 25:75.



**Figure 3c: Skew Ratio 3.** All four samples have  $X_i$  assignment on  $X_m$ . The resulting skew ratio is 0:100. This is a rare occurrence in nature, and is often referred to as non-random assignment or fully skewed [3]

**%Xi or tau ( $\tau$ ):** The ratio of a gene's expression on  $X_i$  over its total expression across both  $X_a$  and  $X_i$ , referred to as %Xi [4] or  $\tau$  [5]. In allele specific expression (ASE) based methods for escape inference,  $\tau$  is compared to the skew ratio. Significant deviation of  $\tau$  from the skew ratio reflects biallelic expression and classifies a gene as escape.

$N$  = expression level

$j$  = gene

$i$  = inactive  $x$  chromosome

$a$  = active  $x$  chromosome

$$\tau = \frac{N_{ij}}{N_{aj} + N_{ij}}$$

*A note about skew: For ASE based methods, data sets with a higher skew ratio (such as 25:75) are strengthened for estimating escape expression due to the lower  $X_i$  expression levels required to infer significance.*

**Tau+ ( $\tau+$ ):** The  $\tau$  (or %Xi) values from the subset of XCI calls which were inferred from samples which were sufficiently skewed (>25:75). This subset is a more robust estimate of ASE and is strengthened for escape calling. [4]

[1] LYON MF. Gene action in the X-chromosome of the mouse (*Mus musculus* L.). *Nature*. 1961;190:372-373. doi:10.1038/190372a0

[2] Carrel L, Brown CJ. When the Lyon(ized chromosome) roars: ongoing expression from an inactive X chromosome. *Philos Trans R Soc Lond B Biol Sci*. 2017;372(1733):20160355. doi:10.1098/rstb.2016.0355

[3] Tukiainen T, Villani AC, Yen A, et al. Landscape of X chromosome inactivation across human tissues [published correction appears in *Nature*. 2018 Mar 7;555(7695):274]. *Nature*. 2017;550(7675):244-248. doi:10.1038/nature24265

[4] Cotton, A.M., Ge, B., Light, N. et al. Analysis of expressed SNPs identifies variable extents of expression from the human inactive X chromosome. *Genome Biol* 14, R122 (2013). doi:10.1186/gb-2013-14-11-r122

[5] Sauteraud Renan, Stahl Jill, James Jesica, Englebright Marisa, Chen Fang, Xiaowei Zhan, Carrel Laura, Liu Dajiang J., Inferring Genes that Escape X Chromosome Inactivation (with XCIR) Reveals Important Contribution of Variable Escape Genes to Sex-biased Diseases. *Genome Research*, accepted.