



MECP2 iPSC Collection at Coriell Institute

| Protein variant | DNA variant | Clonally isolated isogenic control | Sex | Source |
|--------------------------|------------------------|------------------------------------|-----|------------------------|
| Point Mutations | | | | |
| p.R106W | c.C316T | Yes | F | lymphocyte |
| p.R133C | c.C397T | Yes | F | lymphocyte |
| p.R133H | c.G398A | Yes | F | lymphocyte |
| p.T158M | c.C473T | Yes | F | lymphocyte |
| p.R168X | c.C502T | Yes | F | lymphocyte |
| p.R255X | c.C763T | Yes | F | lymphocyte |
| p.R270X | c.C808T | Yes | F | lymphocyte |
| p.R294X | c.C880T | Yes | F | fibroblast |
| p.R306C | c.C916T | Yes | F | lymphocyte |
| Splicing Mutation | | | | |
| --- | c.62+1delGT (intron 1) | Yes | F | Coriell fibro. GM25456 |

MECP2 Fibroblast Collection at Sampled

| Protein variant | DNA variant | Sex | Source |
|---------------------------------------|--|-----|-------------|
| Point Mutations | | | |
| p.P72L | c.C215T | F | skin biopsy |
| p.R106W | c.C316T | F | skin biopsy |
| p.R133C | c.C397T | F | skin biopsy |
| p.T158M | c.C473T | F | skin biopsy |
| p.R168X | c.C502T | F | skin biopsy |
| p.R255X | c.C763T | F | skin biopsy |
| p.R270X | c.C808T | F | skin biopsy |
| p.R294X | c.C880T | F | skin biopsy |
| p.R306C | c.C916T | F | skin biopsy |
| Single Base Deletion Mutations | | | |
| p.P72Rfs*53 | c.215del | M | skin biopsy |
| p.G252Afs | c.753delC | F | skin biopsy |
| p.G269Afs*20 | c.806delG | M/F | skin biopsy |
| Duplicated Region Mutation | | | |
| p.R20EfsX29 | c.44_57dup14 | F | skin biopsy |
| Splicing Mutations | | | |
| --- | c.62+2T>G; IVS1+2T>G in intron 1 | M | skin biopsy |
| --- | IVS3-2A>G | F | skin biopsy |
| Large Deletion Mutations | | | |
| --- | exon 4 partial deletion: codon 377 into 3' UTR | F | skin biopsy |
| --- | 97kb interstitial deletion of XQ28-Q28 | F | skin biopsy |