| Gene Accession No. | Exon | Mutation | Nucleotide change | Amino acid change | Mutation type | VAF in myeloid blasts | VAF in PDCs | VAF in promonocyte | VAF in BM |
|------------------|------|----------|------------------|------------------|---------------|-----------------------|------------|-------------------|---------|
| DNHD1 NM_144666  | 11   | c.T2145C | p.H715H | Silent | Mutation type | 47.57 | 48.55 | 38.91 | |
| LOC440243 NM_001271664 | 15 | c.C1717A | p.R573R | Silent | Mutation type | 0.00 | 0.00 | 15.38 | |
| NOMO1 NM_014287 | 16 | c.A3216G | p.T1072T | Silent | Mutation type | 0.00 | 0.00 | 7.89 | |
| NXPH1 NM_152745 | 7 | c.C567A | p.S189S | Silent | Mutation type | 50.38 | 52.31 | 47.83 | |
| PCDHB10 NM_018930 | 5 | c.C1797T | p.N599N | Silent | Mutation type | 0.00 | 22.22 | 10.30 | |
| PCDHB10 NM_018930 | 5 | c.C1920G | p.L640L | Silent | Mutation type | 9.09 | 0.00 | 6.40 | |
| POTEC NM_001137671 | 18 | c.G492T | p.T164T | Silent | Mutation type | 6.41 | 0.00 | 7.50 | |
| SPEG NM_005876 | 2 | c.G3273A | p.P1091P | Silent | Mutation type | 6.91 | 31.82 | 14.21 | |
| VNN1 NM_004666 | 6 | c.C693A | p.T231T | Silent | Mutation type | 40.40 | 46.82 | 41.80 | |
| ADAM19 NM_033274 | 5 | c.C2564T | p.P855L | Missense Mutation | Mutation type | 48.25 | 45.88 | 37.30 | |
| C4BPA NM_000715 | 1 | c.C358T | p.R120C | Missense Mutation | Mutation type | 54.24 | 46.85 | 40.00 | |
| CAMSAP1 NM_015447 | 9 | c.G2843A | p.G948E | Missense Mutation | Mutation type | 45.31 | 43.27 | 35.44 | |
| CYP21A2 NM_022462 | 6 | c.G1361C | p.R454P | Missense Mutation | Mutation type | 0.00 | 0.00 | 5.13 | |
| HDHD5 NM_033070 | 22 | c.T52G | p.W18G | Missense Mutation | Mutation type | 16.67 | 9.09 | 12.00 | |
| HIF3A NM_022462 | 19 | c.C1483G | p.P495A | Missense Mutation | Mutation type | 59.18 | 55.71 | 35.34 | |
| LRP1B NM_018557 | 2 | c.G11833A | p.G3945R | Missense Mutation | Mutation type | 37.50 | 44.96 | 30.30 | |
| LYVE1 NM_006691 | 11 | c.T698G | p.F233C | Missense Mutation | Mutation type | 0.00 | 0.00 | 17.65 | |
| MUC7 NM_152291 | 4 | c.T779A | p.V260D | Missense Mutation | Mutation type | 11.90 | 0.00 | 5.61 | |
| NPIPA5 NM_001277325 | 16 | c.T770C | p.I257T | Missense Mutation | Mutation type | 5.49 | 0.00 | 6.38 | |
| NPIPA5 NM_001277325 | 16 | c.T659C | p.V220A | Missense Mutation | Mutation type | 7.84 | 0.00 | 8.00 | |
| RBM20 NM_001134363 | 10 | c.C1145T | p.A382V | Missense Mutation | Mutation type | 49.38 | 47.97 | 37.84 | |
| RUNX1 NM_001001890 | 21 | c.G237C | p.W79C | Missense Mutation | Mutation type | 59.09 | 55.32 | 40.20 | |
| SECTM1 NM_003004 | 17 | c.C614T | p.A205V | Missense Mutation | Mutation type | 39.13 | 57.41 | 25.00 | |
| SLC35G4 NM_001282300 | 18 | c.G772A | p.A258T | Missense Mutation | Mutation type | 13.64 | 0.00 | 5.31 | |
| SLCO4A1 NM_016354 | 20 | c.A1882C | p.I628L | Missense Mutation | Mutation type | 0.00 | 0.00 | 10.71 | |
| SPHKAP NM_001142644 | 2 | c.G3536T | p.S1179I | Missense Mutation | Mutation type | 49.27 | 50.00 | 33.33 | |
| SPICE1 NM_144718 | 3 | c.G1980T | p.E660D | Missense Mutation | Mutation type | 48.00 | 46.43 | 25.56 | |
| WASHC1 NM_182905 | 9 | c.G1127A | p.G376D | Missense Mutation | Mutation type | 0.00 | 28.57 | 12.33 | |
| BCOR NM_001123383 | X | c.761delG | p.G254fs frameshift deletion | Missense Mutation | Mutation type | 99.70 | 100.00 | 83.75 | |
| BCORL1 NM_021946 | X | c.514_525del | p.172_175del nonframeshift deletion | Missense Mutation | Mutation type | 97.70 | 100.00 | 71.95 | |
| CHD5 NM_015557 | 1 | c.T5004C | p.D1668D | Silent | Mutation type | 52.50 | 58.93 | 59.40 | 50.00 |