

SUPPLEMENTARY METHODS

Detailed PCR protocol for amplification of genomic DNA and cDNA, isolated from PMBL samples, followed by sequencing.

PCR reaction was prepared using 0,2 mM dNTP, 0,5 pmol of each primer (suppl table 1), 2,5 U Taq Polymerase (GE Helthcare, Little Chalfont, UK) and either 50 ng of genomic DNA or 2 µl of cDNA, in a final 50 µl volume. The following touchdown PCR program was used: denaturation at 95°C for 5min; 2 cycles of denaturation at 95°C for 1 min, annealing at 62°C for 1min, elongation at 72°C for 1 min; 2 cycles of denaturation at 95°C for 1 min, annealing at 60°C for 1 min, elongation at 72°C for 1 min; 2 cycles denaturation at 95°C for 1 min, annealing at 58°C for 1 min, elongation at 72°C for 1 min; 30 cycles of denaturation at 95°C for 1 min, annealing at 56°C for 1 min, elongation 72°C for 1 min; and final elongation step at 72°C for 10 min.

Oligonucleotides used for PCR amplification and quantification or sequencing.

| primers | matrix | technique | 5' sequence 3' |
|----------------|--------------|------------------|---------------------------|
| STAT6 cDNA for | cDNA | PCR and sequence | GTGCTCTTCTCTGCCAGCTT |
| STAT6 cDNA rev | cDNA | PCR and sequence | ATCTGTGGAGAGCCATCCTG |
| STAT6 ex12 for | gDNA | PCR and sequence | CCTGTCCTCACCTCTTCAG |
| STAT6 ex12 rev | gDNA | PCR and sequence | CAGGCCCATGAGAAAGTGTT |
| STAT6 ex13 for | gDNA | PCR and sequence | GCCTGCAGTGCTCTTCTT |
| STAT6 ex13 rev | gDNA | PCR and sequence | CCAGGGATGAAGAGCTTGG |
| STAT6 ex14 for | gDNA | PCR and sequence | CTCACACCTTTCCCCCTCTC |
| STAT6 ex14 rev | gDNA | PCR and sequence | CTGGTGTATGGCTGCTCAGA |
| SOCS1 for | cDNA or gDNA | PCR and sequence | ATGGTAGCACACAACCAGGTGG |
| SOCS1 rev | cDNA or gDNA | PCR and sequence | TCAAATCTGGAAGGGGAAGGAGCTC |
| JAK2 for | gDNA | Real time PCR | AAGCCACTGCCAGAACTTG |
| JAK2 rev | gDNA | Real time PCR | ACTGAATTCCACCGTTTCCA |
| B2M for | gDNA | Real time PCR | TGTCTTTCAGCAAGGACTGG |
| B2M rev | gDNA | Real time PCR | GATGCTGCTTACATGTCTCG |