| Motif | Group Specificity | Group Specificity Rank | Site Specificity | Site Specificy Rank | Testset Sites | Average Testset Sites | StDev Testset Sites | TRANSFAC Correlation | TRANSFAC ID | Motif Finders | Sets | Repeats | Logo |
|---|---|---|---|---|---|---|---|---|---|---|---|---|---|
| 1 | 5.70e-015 | 2 | 8.80e-014 | 3 | 7 | 0.64 | 0.50 | 0.69 | HAND1E | AA, BP, MD | 1, 2, 3, 4, 5 | TR | |
| 2 | 2.90e-012 | 5 | 3.40e-010 | 14 | 10 | 0.91 | 0.54 | 0.62 | TATA | AA | 1, 2, 3, 4, 5 | TR | |
| 3 | 2.10e-011 | 10 | 1.90e-009 | 17 | 14 | 1.30 | 0.90 | 0.70 | POU1F1 | AA, BP | 1, 2, 3, 4, 5 | | |
| 4 | 3.40e-011 | 12 | 9.00e-011 | 8 | 8 | 0.73 | 0.90 | 0.62 | ICSBP | AA | 1, 2, 3, 4, 5 | PAL,TR | |
| 5 | 3.20e-010 | 18 | 1.30e-007 | 32 | 14 | 1.30 | 1.00 | 0.69 | ELF1 | AA, ME | 1, 2, 3, 4, 5 | | |
| 6 | 3.30e-010 | 19 | 7.10e-010 | 15 | 5 | 0.45 | 0.69 | 0.74 | NFY | AA, BP, ME | 1, 2, 3, 4, 5 | | |
| 7 | 1.00e-009 | 22 | 1.20e-008 | 22 | 9 | 0.82 | 0.87 | 0.73 | IRF1 | AA | 1, 2, 3, 4, 5 | TR | |
| 8 | 2.40e-008 | 36 | 1.30e-007 | 32 | 12 | 1.10 | 1.50 | 0.64 | NFAT_Q | AA | 1, 2, 3, 4, 5 | PAL,TR | |
| 9 | 2.70e-008 | 37 | 5.60e-008 | 26 | 6 | 0.55 | 0.93 | 0.64 | ETS2 | AA, ME | 1, 2, 3, 4, 5 | | |
| 10 | 2.10e-007 | 40 | 1.20e-006 | 39 | 7 | 0.64 | 0.67 | 0.67 | SMAD3 | BP | 3 | PAL | |
| 11 | 3.00e-007 | 43 | 6.70e-006 | 53 | 11 | 1.00 | 0.77 | 0.75 | AML | BP, MD, ME | 1, 2, 3, 4, 5 | | |
| 12 | 3.10e-007 | 44 | 6.60e-006 | 52 | 12 | 1.10 | 0.70 | 0.66 | SP1_Q2 | BP, ME | 1, 2, 4, 5 | TR | |
| 13 | 3.60e-007 | 45 | 1.90e-005 | 61 | 12 | 1.10 | 0.83 | 0.71 | NFAT | BP, ME | 2, 3 | | |
| 14 | 3.80e-007 | 48 | 2.20e-006 | 43 | 8 | 0.73 | 0.79 | 0.65 | P300 | BP | 5 | | |
| 15 | 5.50e-007 | 52 | 1.70e-006 | 40 | 9 | 0.82 | 1.30 | 0.66 | CACBIN | AA, BP | 1, 2, 3, 4, 5 | TR | |
| 16 | 5.50e-007 | 52 | 1.20e-005 | 55 | 11 | 1.00 | 0.77 | 0.65 | PXR | BP, ME | 1, 2, 3, 4, 5 | PAL | |
| 17 | 7.00e-007 | 56 | 2.20e-006 | 43 | 11 | 1.00 | 2.00 | 0.61 | CEBPGA | MD, ME | 1, 2, 3, 4, 5 | PAL,TR | |
| 18 | 7.50e-007 | 57 | 9.90e-005 | 76 | 15 | 1.40 | 0.67 | 0.66 | LHX3 | BP | 2, 3, 4 | PAL | |
| 19 | 8.40e-007 | 58 | 1.60e-005 | 59 | 11 | 1.00 | 0.77 | 0.69 | FREAC7 | AA, BP | 1, 2, 3, 4, 5 | | |
| 20 | 2.40e-006 | 61 | 1.30e-005 | 56 | 9 | 0.82 | 0.87 | 0.67 | MEF3 | BP | 1, 3, 4, 5 | | |
| 21 | 3.10e-006 | 62 | 5.70e-005 | 71 | 13 | 1.20 | 1.10 | 0.62 | FREAC7 | AA, BP, ME | 1, 2, 3, 5 | | |
| 22 | 5.60e-006 | 67 | 2.50e-005 | 64 | 12 | 1.10 | 1.40 | 0.62 | ELF1 | AA, BP | 1, 2, 3, 4, 5 | | |
| 23 | 6.00e-006 | 68 | 1.80e-005 | 60 | 7 | 0.64 | 0.81 | 0.60 | TFIII | AA, ME | 1, 4 | | |
| 24 | 7.30e-006 | 69 | 1.10e-004 | 78 | 11 | 1.00 | 0.89 | 0.63 | PR | BP | 1, 2, 3, 4, 5 | | |
| 25 | 7.40e-006 | 70 | 2.30e-005 | 63 | 7 | 0.64 | 0.81 | 0.62 | GR_Q6 | BP, ME | 4, 5 | | |
| 26 | 9.90e-006 | 74 | 4.80e-005 | 69 | 8 | 0.73 | 0.65 | 0.68 | SMAD4 | BP, ME | 1, 2, 3, 4 | | |
| 27 | 1.30e-005 | 76 | 3.70e-005 | 66 | 11 | 1.00 | 2.00 | 0.65 | SRF | MD | 1, 2, 3, 4, 5 | PAL,TR | |
| 28 | 2.10e-005 | 78 | 3.20e-004 | 83 | 12 | 1.10 | 0.94 | 0.76 | TATA | ME | 1, 2, 3, 4, 5 | | |
| 29 | 2.30e-005 | 80 | 6.40e-005 | 72 | 8 | 0.73 | 0.90 | 0.62 | IRF | MD | 1, 2, 3, 4, 5 | TR | |
| 30 | 3.10e-005 | 82 | 1.60e-003 | 95 | 19 | 1.70 | 1.00 | 0.63 | MYB_Q5 | BP | 1, 2 | | |
| 31 | 4.50e-005 | 85 | 3.50e-004 | 84 | 11 | 1.00 | 1.10 | 0.67 | HOXA4 | AA, BP, ME | 1, 2, 3, 4 | PAL | |
| 32 | 9.90e-005 | 92 | 1.10e-003 | 91 | 13 | 1.20 | 0.98 | 0.62 | CHOP | BP | 2, 4 | PAL | |