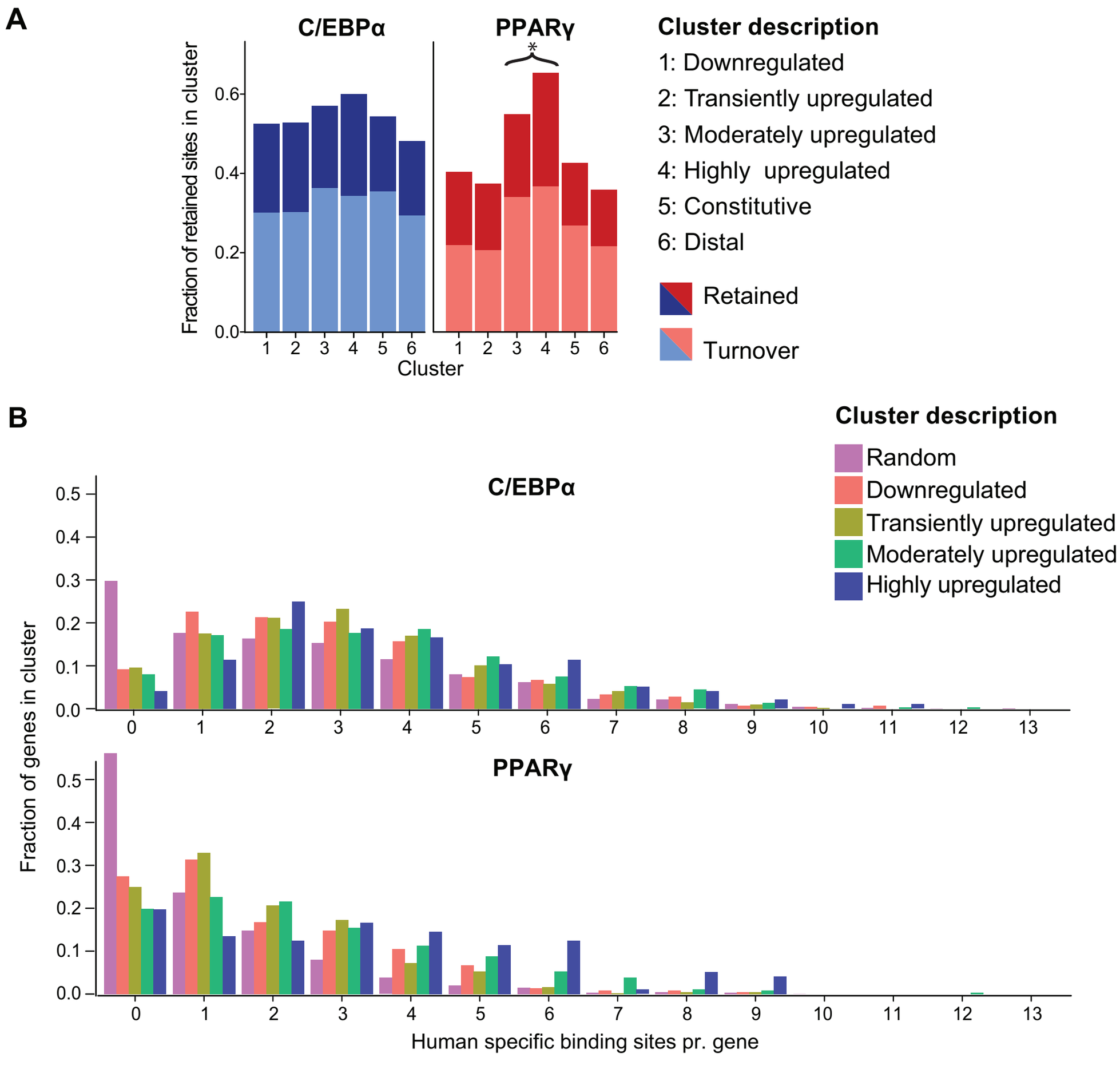
A
C/EBPα
PPARγ
*
Fraction of retained sites in cluster
0.0
0.2
0.4
0.6
1 2 3 4 5 6
1 2 3 4 5 6
Cluster
Cluster description
1: Downregulated
2: Transiently upregulated
3: Moderately upregulated
4: Highly upregulated
5: Constitutive
6: Distal
Retained
Turnover
B
Cluster description
Random
Downregulated
Transiently upregulated
Moderately upregulated
Highly upregulated
C/EBPα
PPARγ
Fraction of genes in cluster
0.0
0.1
0.2
0.3
0.4
0.5
0 1 2 3 4 5 6 7 8 9 10 11 12 13
Human specific binding sites pr. gene