							group	1.5	group	
							ITGB8_M1	1.5	00)4)4–185–KD
							FN1_M1		Modu	
							IL32_M1	0		1 immune system process
							ETS1_M1	-0.5		2 cytokine production 3 response to cytokine
							CYP26B1_M1	-1		4 PI3K–Akt signaling pathway 5 regulation of MAPK cascade
							ITGA1_M1	-1.5	M	6 nervous system development
							EPAS1_M1		M [*]	7 regulation of RNA metabolic process
							FN1_M2			
							IL32_M2			
							IGF2BP1_M2			
							CPNE1_M3			
							ERBB3_M4			
							ITGB8_M4			
							FN1_M4			
							ITGA1_M4			
							FN1_M5			
							FOXO1_M5			
							ITGA1_M5			
							ID3_M6			
							ERBB3_M6			
ľ							PCDHGB4_M6			
ı							PCDHGA8_M6			
							PCDHGA5_M6			
							FN1_M6			
							PCDHGA6_M6			
							PCDHGB3_M6			
							ANXA2_M6			
							TBX3_M6			
							CPNE1_M6			
							IGF2BP1_M6			
							VCL_M6			
							ZIC2_M6			
							PCDHGB1_M6			
							ITGA1_M6			
							ID3_M7			
							ETS1_M7			
							ANXA2_M7			
							TBX3_M7			
							PPP1R13L_M7			
							IGF2BP1_M7			
							KANK2_M7			
							ZIC2_M7			
							FOXO1_M7			
							EPAS1_M7			
					_		RBM20_M7			
Module	004-A	004-B	004-C	004–185–KD–A	004–185–KD–B	004-185-KD-C				
ile	Þ	w	O	185-	185-	185-k				
				⟨D-A	⟨D-B	⟨D-C				