

**A**

Top molecules	Exp. <i>P</i> Value	<i>HUGO</i> gene name: function
BCR	9,39E-12	breakpoint cluster region: not clear
MIF	3,99E-09	macrophage migration inhibitory factor: suppression of anti-inflammatory effects
COMT	4,06E-08	cartilage oligomeric matrix protein: noncollagenous extracellular matrix (ECM) protein
PRPF31	3,65E-07	pre-mRNA processing factor 31 homolog: spliceosome complex activation
MCM5*	9,94E-07	minichromosome maintenance complex component 5: initiation of DNA replication
CDC45	2,44E-06	cell division cycle 45 homolog: initiation of DNA replication
CDC42EP1	3,05E-06	CDC42 effector protein (Rho GTPase binding) 1: actin cytoskeleton reorganization
CNOT3	3,07E-06	CCR4-NOT transcription complex, subunit 3: general transcription regulation complex
UBE2S	1,10E-05	ubiquitin-conjugating enzyme E2S:ubiquitin activating enzyme

**B**

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