

S\_PAM3CSK4\_4H\_BMDC\_DN, GSE17721\_LPS\_VS\_PAM3CSK4\_4H\_BMDC\_DN

GSE17721\_4\_VS\_24H\_GARDIQUIMOD\_BMDC\_DN, GSE17721\_4\_VS\_24H\_GARDIQUIMOD\_BMDC\_DN  
GSE46606\_UNSTIM\_VS\_CD40L\_IL2\_IL5\_3DAY\_STIMULATED\_IRF4MID\_SORTED\_BCELL\_UP, GSE46606\_UNSTIM\_VS\_CD40L\_IL2\_IL5\_3DAY\_STIMULATED\_IRF4MID\_SORTED\_BCELL\_UP  
GSE17721\_PAM3CSK4\_VS\_CPG\_6H\_BMDC\_UP, GSE17721\_PAM3CSK4\_VS\_CPG\_6H\_BMDC\_UP  
GSE17721\_0.5H\_VS\_12H\_PAM3CSK4\_BMDC\_DN, GSE17721\_0.5H\_VS\_12H\_PAM3CSK4\_BMDC\_DN  
GSE17721\_LPS\_VS\_GARDIQUIMOD\_16H\_BMDC\_DN, GSE17721\_LPS\_VS\_GARDIQUIMOD\_16H\_BMDC\_DN  
REACTOME\_METABOLISM\_OF\_VITAMINS\_AND\_COFACTORS, REACTOME\_METABOLISM\_OF\_VITAMINS\_AND\_COFACTORS  
GOBP\_PROTEIN\_N\_LINKED\_GLYCOSYLATION, GOBP\_PROTEIN\_N\_LINKED\_GLYCOSYLATION  
REACTOME\_METABOLISM\_OF\_WATER\_SOLUBLE\_VITAMINS\_AND\_COFACTORS, REACTOME\_METABOLISM\_OF\_WATER\_SOLUBLE\_VITAMINS\_AND\_COFACTORS  
MIR6860, MIR6860  
GARGALOVIC\_RESPONSE\_TO\_OXIDIZED\_PHOSPHOLIPIDS\_TURQUOISE\_UP, GARGALOVIC\_RESPONSE\_TO\_OXIDIZED\_PHOSPHOLIPIDS\_TURQUOISE\_UP  
GOBP\_WATER\_SOLUBLE\_VITAMIN\_METABOLIC\_PROCESS, GOBP\_WATER\_SOLUBLE\_VITAMIN\_METABOLIC\_PROCESS  
GOBP\_VITAMIN\_METABOLIC\_PROCESS, GOBP\_VITAMIN\_METABOLIC\_PROCESS  
MIR877\_5P, MIR877\_5P  
GOBP\_REGULATION\_OF\_REPRODUCTIVE\_PROCESS, GOBP\_REGULATION\_OF\_REPRODUCTIVE\_PROCESS  
GOBP\_FATTY\_ACYL\_COA\_BIOSYNTHETIC\_PROCESS, GOBP\_FATTY\_ACYL\_COA\_BIOSYNTHETIC\_PROCESS  
GOMF\_TRANSCRIPTION\_COREPRESSOR\_BINDING, GOMF\_TRANSCRIPTION\_COREPRESSOR\_BINDING  
MIR3170, MIR3170  
VALK\_AML\_CLUSTER\_7, VALK\_AML\_CLUSTER\_7  
HP\_PROGRESSIVE\_EXTRAPYRAMIDAL\_MOVEMENT\_DISORDER, HP\_PROGRESSIVE\_EXTRAPYRAMIDAL\_MOVEMENT\_DISORDER  
GOBP\_POSITIVE\_REGULATION\_OF\_UBIQUITIN\_PROTEIN\_LIGASE\_ACTIVITY, GOBP\_POSITIVE\_REGULATION\_OF\_UBIQUITIN\_PROTEIN\_LIGASE\_ACTIVITY  
GOMF\_INSULIN\_RECEPTOR\_BINDING, GOMF\_INSULIN\_RECEPTOR\_BINDING  
REACTOME\_INWARDLY\_RECTIFYING\_K\_CHANNELS, REACTOME\_INWARDLY\_RECTIFYING\_K\_CHANNELS  
WP\_OMEGA9\_FA\_SYNTHESIS, WP\_OMEGA9\_FA\_SYNTHESIS  
REACTOME\_SYNTHESIS\_OF\_PROSTAGLANDINS\_PG\_AND\_THROMBOXANES\_TX, REACTOME\_SYNTHESIS\_OF\_PROSTAGLANDINS\_PG\_AND\_THROMBOXANES\_TX  
WP\_PURINERGIC\_SIGNALING, WP\_PURINERGIC\_SIGNALING  
GOMF\_NUCLEOTIDE\_RECEPTOR\_ACTIVITY, GOMF\_NUCLEOTIDE\_RECEPTOR\_ACTIVITY  
MODULE\_367, MODULE\_367  
MCGARVEY\_SILENCED\_BY\_METHYLATION\_IN\_COLON\_CANCER, MCGARVEY\_SILENCED\_BY\_METHYLATION\_IN\_COLON\_CANCER  
MIR4638\_5P, MIR4638\_5P  
REACTOME\_ESTROGEN\_BIOSYNTHESIS, REACTOME\_ESTROGEN\_BIOSYNTHESIS  
GOBP\_PROTEIN\_KINASE\_C\_ACTIVATING\_G\_PROTEIN\_COUPLED\_RECEPTOR\_SIGNALING\_PATHWAY, GOBP\_PROTEIN\_KINASE\_C\_ACTIVATING\_G\_PROTEIN\_COUPLED\_RECEPTOR\_SIGNALING\_PATHWAY  
GOBP\_REGULATION\_OF\_RESPIRATORY\_SYSTEM\_PROCESS, GOBP\_REGULATION\_OF\_RESPIRATORY\_SYSTEM\_PROCESS  
GOMF\_S\_METHYLTRANSFERASE\_ACTIVITY, GOMF\_S\_METHYLTRANSFERASE\_ACTIVITY  
GOMF\_TESTOSTERONE\_17\_BETA\_DEHYDROGENASE\_NADPPLUS\_ACTIVITY, GOMF\_TESTOSTERONE\_17\_BETA\_DEHYDROGENASE\_NADPPLUS\_ACTIVITY  
GOBP\_CELLULAR\_RESPONSE\_TO\_AMMONIUM\_ION, GOBP\_CELLULAR\_RESPONSE\_TO\_AMMONIUM\_ION  
COLE\_BLOOD\_FLUZONE\_FLUARIX\_AGE\_03\_17YO\_7DY\_DN, COLE\_BLOOD\_FLUZONE\_FLUARIX\_AGE\_03\_17YO\_7DY\_DN  
GOBP\_MIDGUT\_DEVELOPMENT, GOBP\_MIDGUT\_DEVELOPMENT