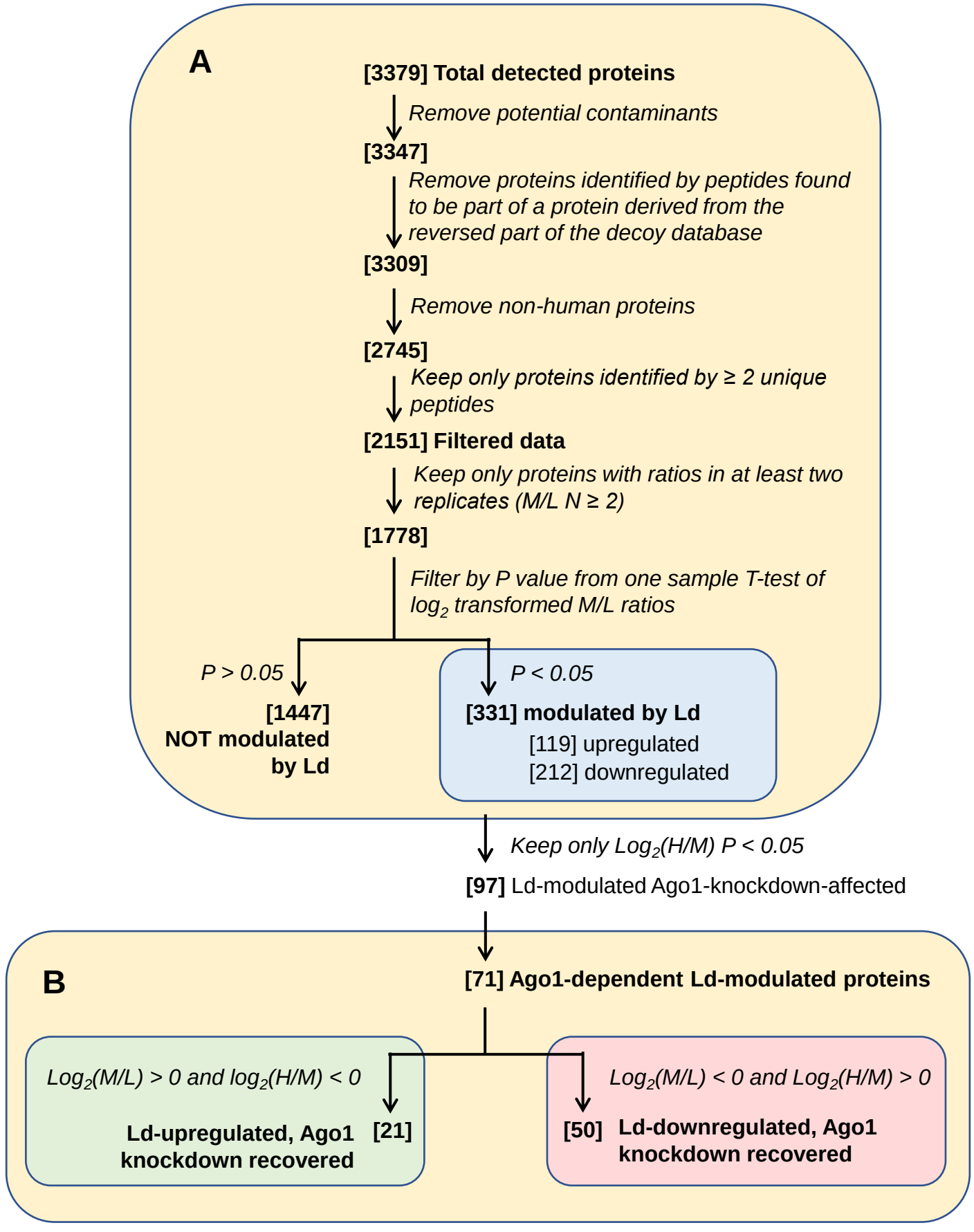
A
[3379] Total detected proteins
Remove potential contaminants
[3347]
Remove proteins identified by peptides found to be part of a protein derived from the reversed part of the decoy database
[3309]
Remove non-human proteins
[2745]
Keep only proteins identified by ≥ 2 unique peptides
[2151] Filtered data
Keep only proteins with ratios in at least two replicates (M/L N ≥ 2)
[1778]
Filter by P value from one sample T-test of log2 transformed M/L ratios
P > 0.05
[1447] NOT modulated by Ld
P < 0.05
[331] modulated by Ld
[119] upregulated
[212] downregulated
Keep only Log2(H/M) P < 0.05
[97] Ld-modulated Ago1-knockdown-affected
B
[71] Ago1-dependent Ld-modulated proteins
Log2(M/L) > 0 and log2(H/M) < 0
[21] Ld-upregulated, Ago1 knockdown recovered
Log2(M/L) < 0 and Log2(H/M) > 0
[50] Ld-downregulated, Ago1 knockdown recovered