

Figure S1 The most enriched GO terms between HP_C and nHP_C. a, biological process; b, molecular function.

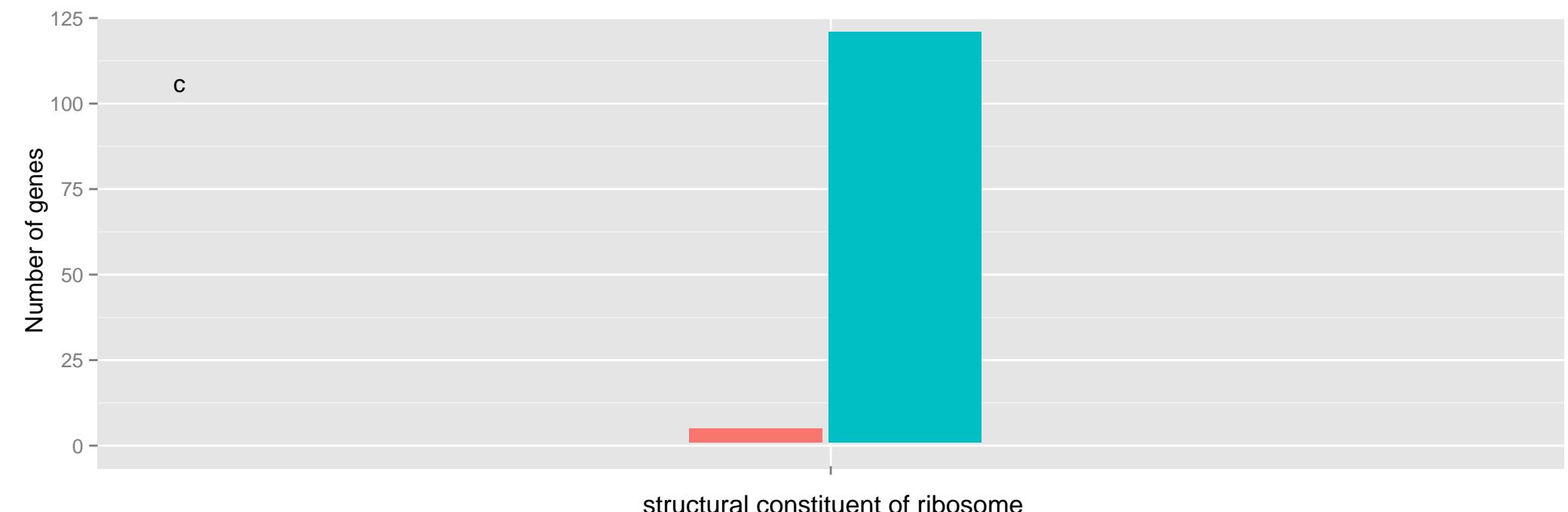
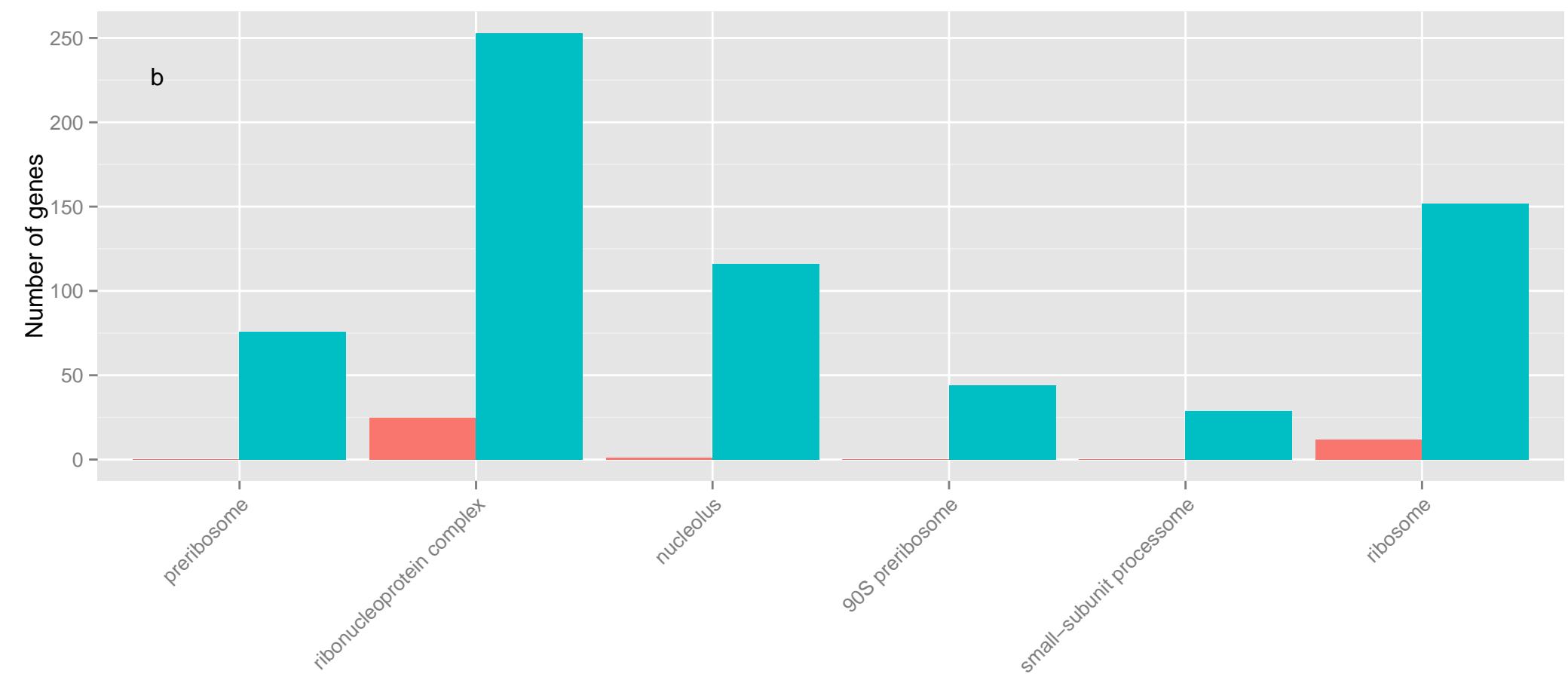
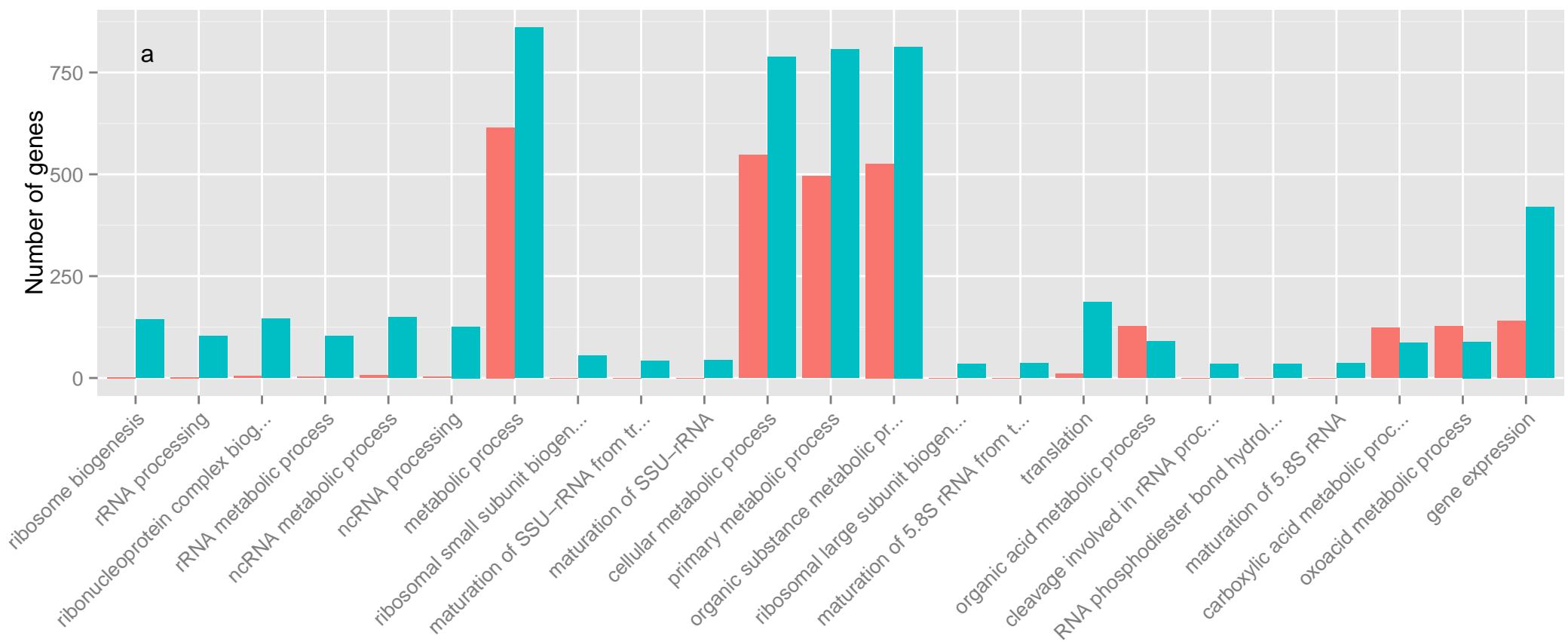


Figure S2 The most enriched GO terms between HP_M and nHP_M. a, biological process; b, cellular component; c, molecular function.

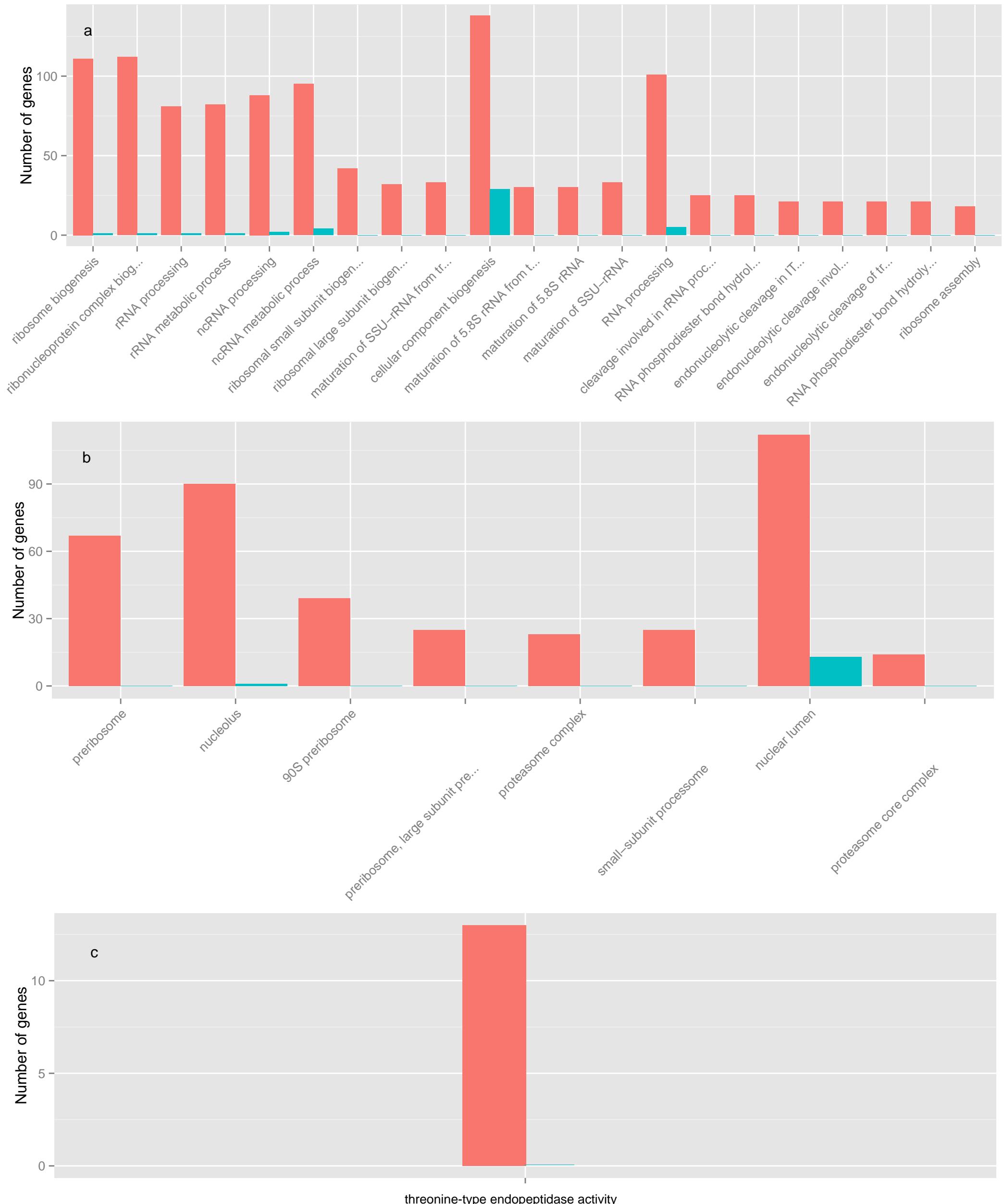
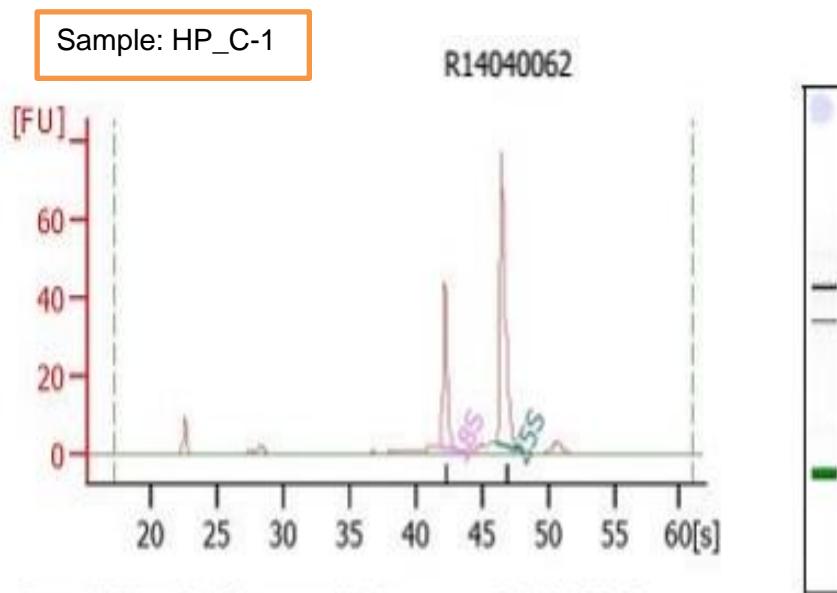


Figure S3 The most enriched GO terms between nHP_M and nHP_C. a, biological process; b, cellular component; c, molecular function.

Figure S5 The quality and the quantity of the RNA samples were checked by using Agilent 2100 Bioanalyzer System, agarose gel electrophoresis and NanoDrop Lite Spectrophotometer.

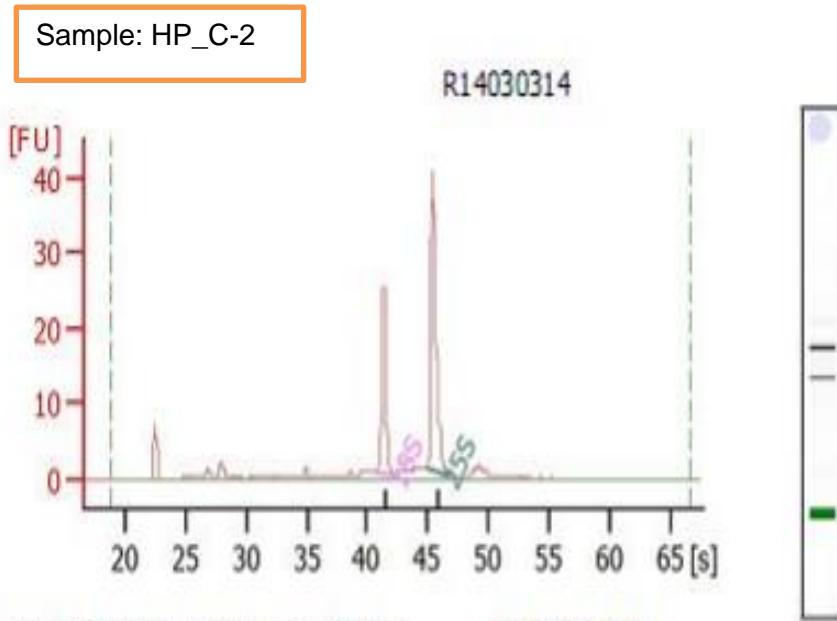


Overall Results for sample 9 : R14040062

RNA Area: 181.6
 RNA Concentration: 163 ng/ μ l
 rRNA Ratio [25s / 18s]: 2.0
 RNA Integrity Number (RIN): 9.5 (B.02.08)
 Result Flagging Color:
 Result Flagging Label: RIN: 9.50

Fragment table for sample 9 : R14040062

Name	Start Time [s]	End Time [s]	Area	% of total Area
18S	41.68	43.23	37.6	20.7
25S	46.06	48.01	76.2	42.0



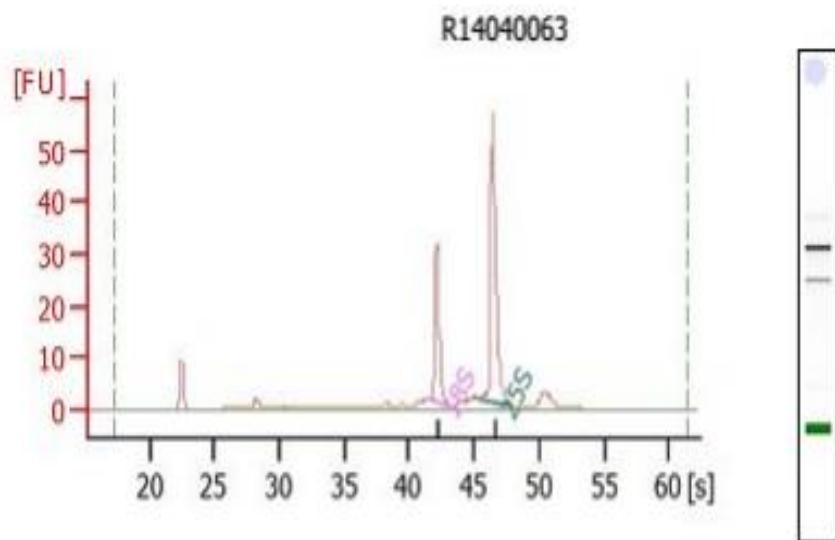
Overall Results for sample 6 : R14030314

RNA Area: 112.4
 RNA Concentration: 69 ng/ μ l
 rRNA Ratio [25s / 18s]: 2.1
 RNA Integrity Number (RIN): 9.4 (B.02.08)
 Result Flagging Color:
 Result Flagging Label: RIN: 9.40

Fragment table for sample 6 : R14030314

Name	Start Time [s]	End Time [s]	Area	% of total Area
18S	40.69	42.34	20.7	18.4
25S	44.95	46.89	43.2	38.4

Sample: HP_M-1



Overall Results for sample 10 : R14040063

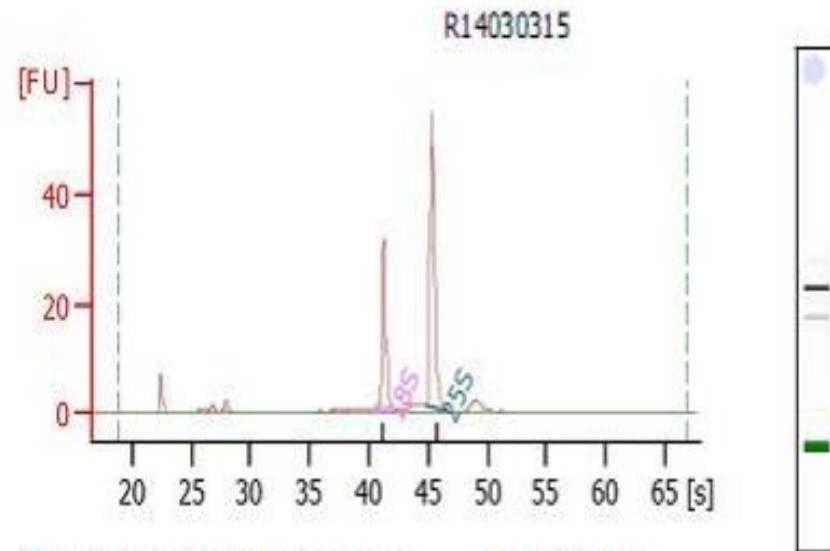
RNA Area: 150.7
RNA Concentration: 135 ng/ μ l
rRNA Ratio [25s / 18s]: 2.1
RNA Integrity Number (RIN): 9.1 (B.02.08)
Result Flagging Color:

Result Flagging Label: RIN: 9.10

Fragment table for sample 10 : R14040063

Name	Start Time [s]	End Time [s]	Area	% of total Area
18S	41.61	43.08	27.3	18.1
25S	45.45	48.03	56.3	37.4

Sample: HP_M-2



Overall Results for sample 7 : R14030315

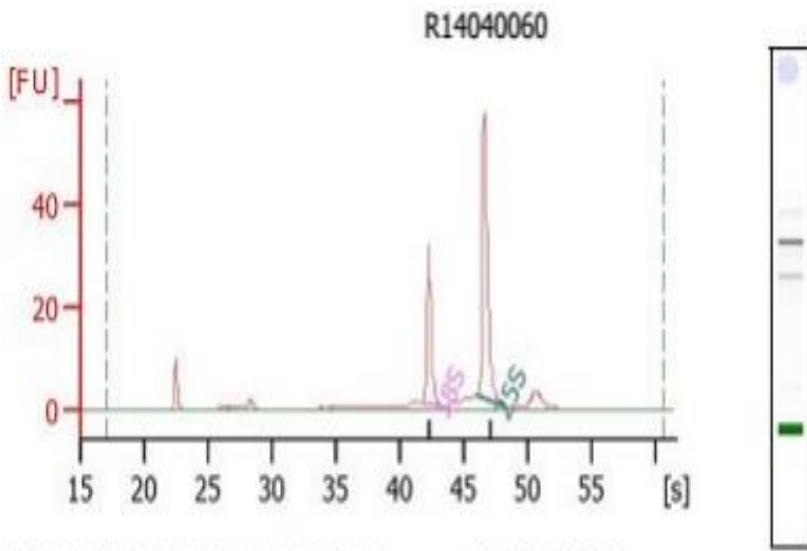
RNA Area: 126.2
RNA Concentration: 77 ng/ μ l
rRNA Ratio [25s / 18s]: 2.1
RNA Integrity Number (RIN): 9.6 (B.02.08)
Result Flagging Color:

Result Flagging Label: RIN: 9.60

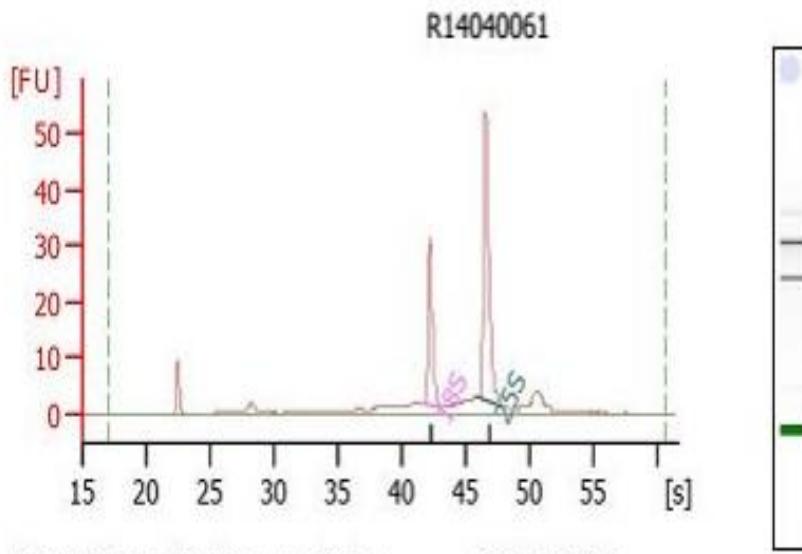
Fragment table for sample 7 : R14030315

Name	Start Time [s]	End Time [s]	Area	% of total Area
18S	40.54	42.04	25.5	20.2
25S	44.76	46.84	54.1	42.9

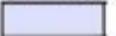
Sample: nHP_C-1



Sample: nHP_C-2



Overall Results for sample 7 : R14040060

RNA Area: 150.2
RNA Concentration: 135 ng/ μ l
rRNA Ratio [25s / 18s]: 2.0
RNA Integrity Number (RIN): 9.2 (B.02.08)
Result Flagging Color: 
Result Flagging Label: RIN: 9.20

Fragment table for sample 7 : R14040060

Name	Start Time [s]	End Time [s]	Area	% of total Area
18S	41.66	43.20	27.4	18.2
25S	46.01	48.21	54.9	36.6

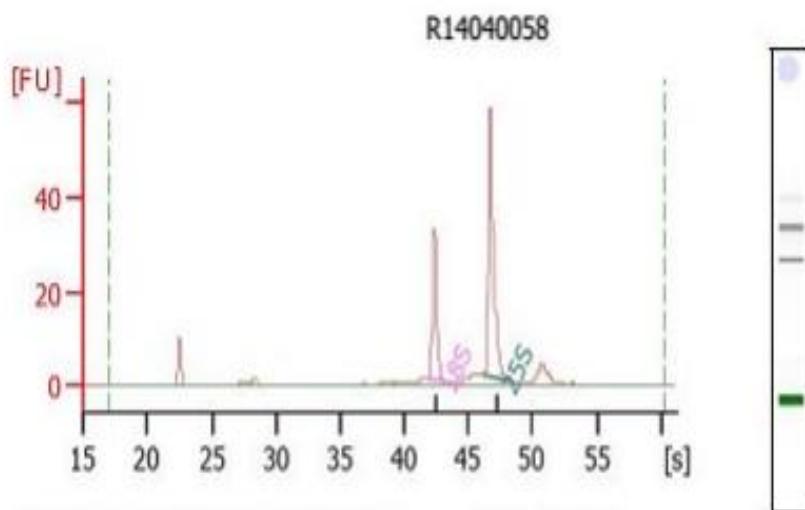
Overall Results for sample 8 : R14040061

RNA Area: 154.5
RNA Concentration: 138 ng/ μ l
rRNA Ratio [25s / 18s]: 2.0
RNA Integrity Number (RIN): 8.9 (B.02.08)
Result Flagging Color: 
Result Flagging Label: RIN: 8.90

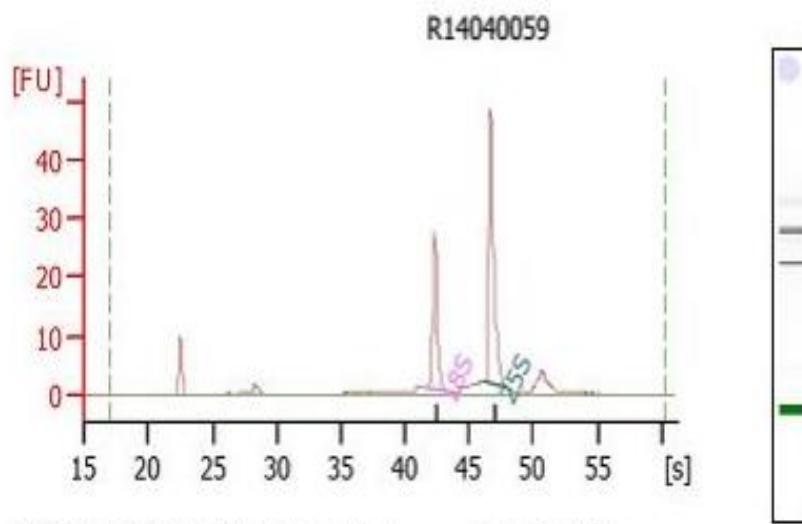
Fragment table for sample 8 : R14040061

Name	Start Time [s]	End Time [s]	Area	% of total Area
18S	41.66	43.15	26.7	17.3
25S	45.92	47.90	52.2	33.8

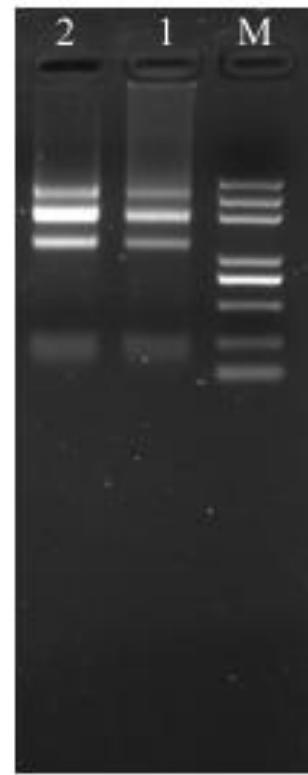
Sample: nHP_M-1



Sample: nHP_M-2

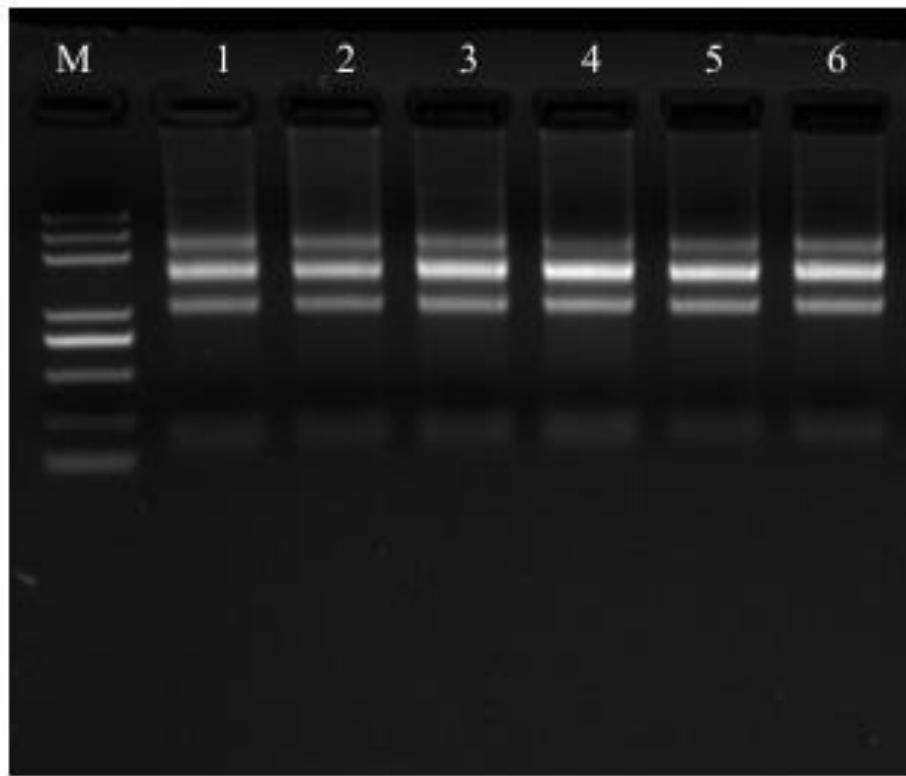


Agarose gel electrophoresis test



Note: Lane M, Trans2KPlus Marker; Lane 1, HP_C-1; Lane 2, HP_M-1.

Agarose gel electrophoresis test



Note: Lane M, Trans2KPlus Marker; Lane 1, nHP_M-1; Lane 2, nHP_M-2; Lane 3, nHP_C-1; ; Lane 4, nHP_C-2; ; Lane 5, HP_C-2; ; Lane 6, HP_M-2.

Sample name	Sample ID	Concentration (ng/µL)	OD260/280	OD260/230	25S/18S	RNA integrity number (RIN)	Inspection
HP_C-1	R14040062	1220	2.279	0.366	2.0	9.5	Pass
HP_C-2	R140303 14	690.00	2.161	0.926	2.1	9.4	Pass
HP_M-1	R14040063	1984	2.238	0.563	2.1	9.1	Pass
HP_M-2	R140303 15	1540.00	2.167	2.172	2.1	9.6	Pass
nHP_M-1	R14040058	2074	1.945	1.087	2.0	9.6	Pass
nHP_M-2	R14040059	1848	1.929	0.857	1.9	9.6	Pass
nHP_C-1	R14040060	2198	1.882	0.646	2.0	9.2	Pass
nHP_C-2	R14040061	1562	2.189	0.263	2.0	8.9	Pass