

Sampling condition	Growth rate (h ⁻¹)	q_{glucose} (mmol.g ⁻¹ .h ⁻¹)	n_{lactate} (mmol.g ⁻¹ .h ⁻¹)
(1) normal	0.88	22 ± 3	42 ± 2
(2) stress : isoleucine depletion	0.05	7 ± 1	15 ± 1

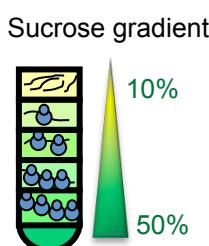


Translation arrest with chloramphenicol and sampling

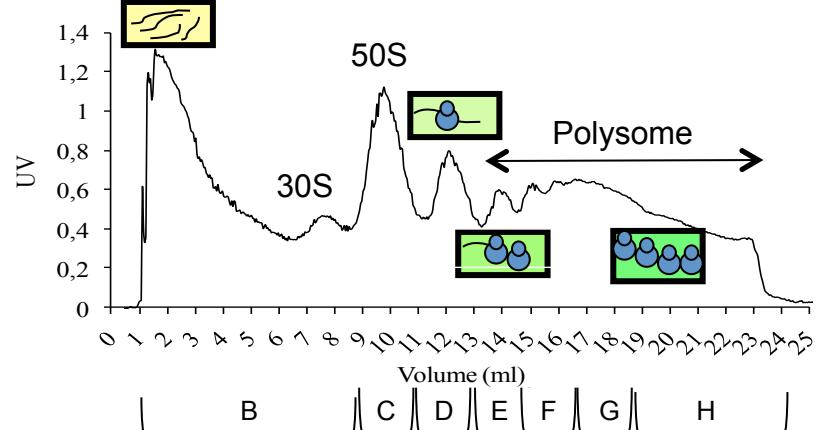


Size separation of ribosome-mRNA complexes according to ribosome number

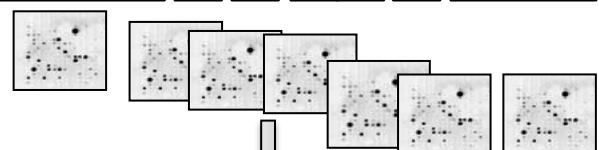
(Example of a polysomal profile in normal condition (1))



Elution



Quantification of mRNA levels in each fraction by microarrays



Statistical data processing

Determination of the number of ribosomes bound to each mRNA molecules

