

A

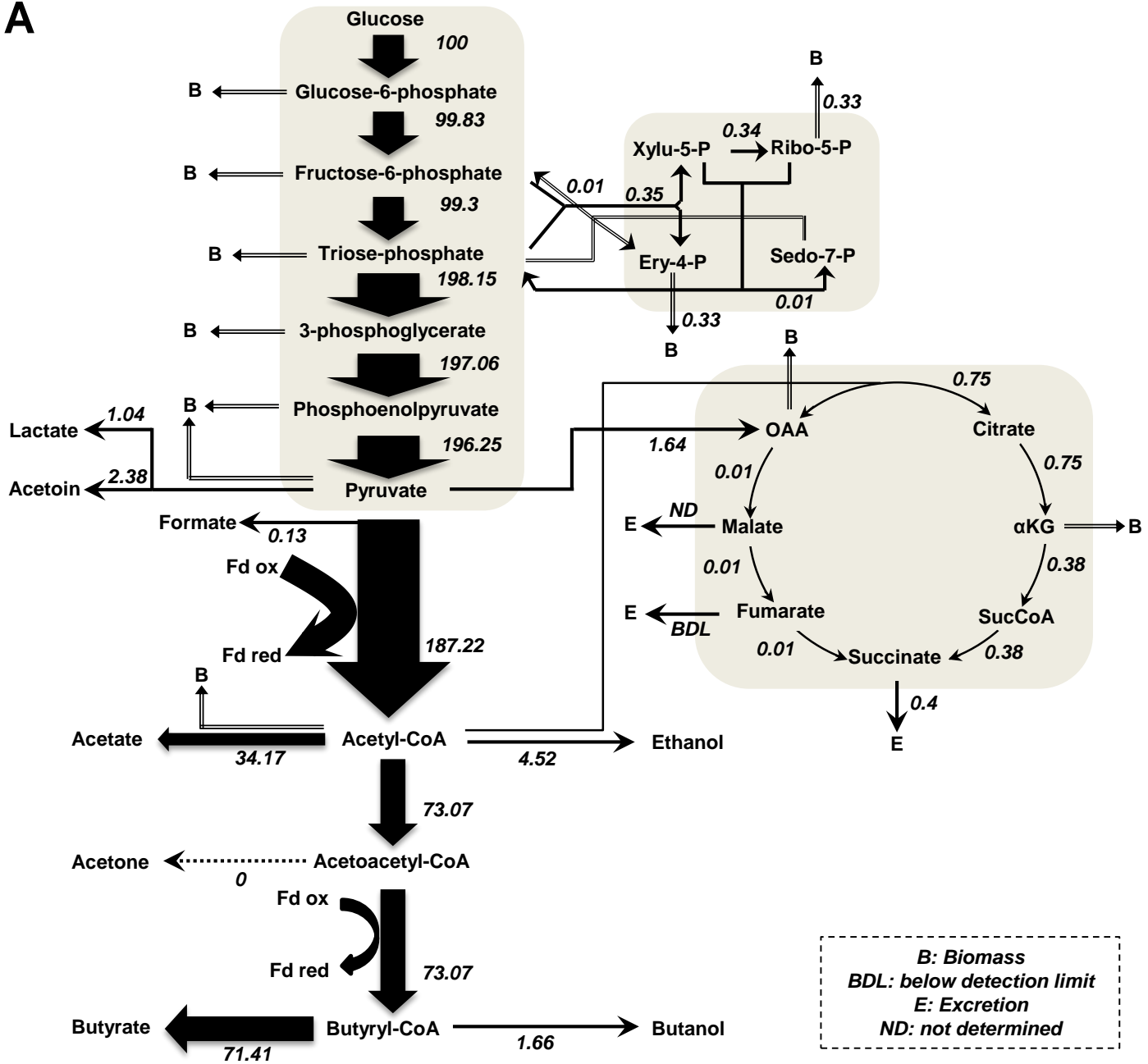


Fig. S3. Metabolic flux map of *C. acetobutylicum* in acidogenesis (A), solventogenesis (B), alcohologenesis (C). All values are normalized to flux of initial carbon source (mmol/gDCW/h). Glucose flux normalized as 100 in the acidogenesis and solventogenesis, and sum of glucose and half of glycerol normalized as 100 in alcohologenesis.

B

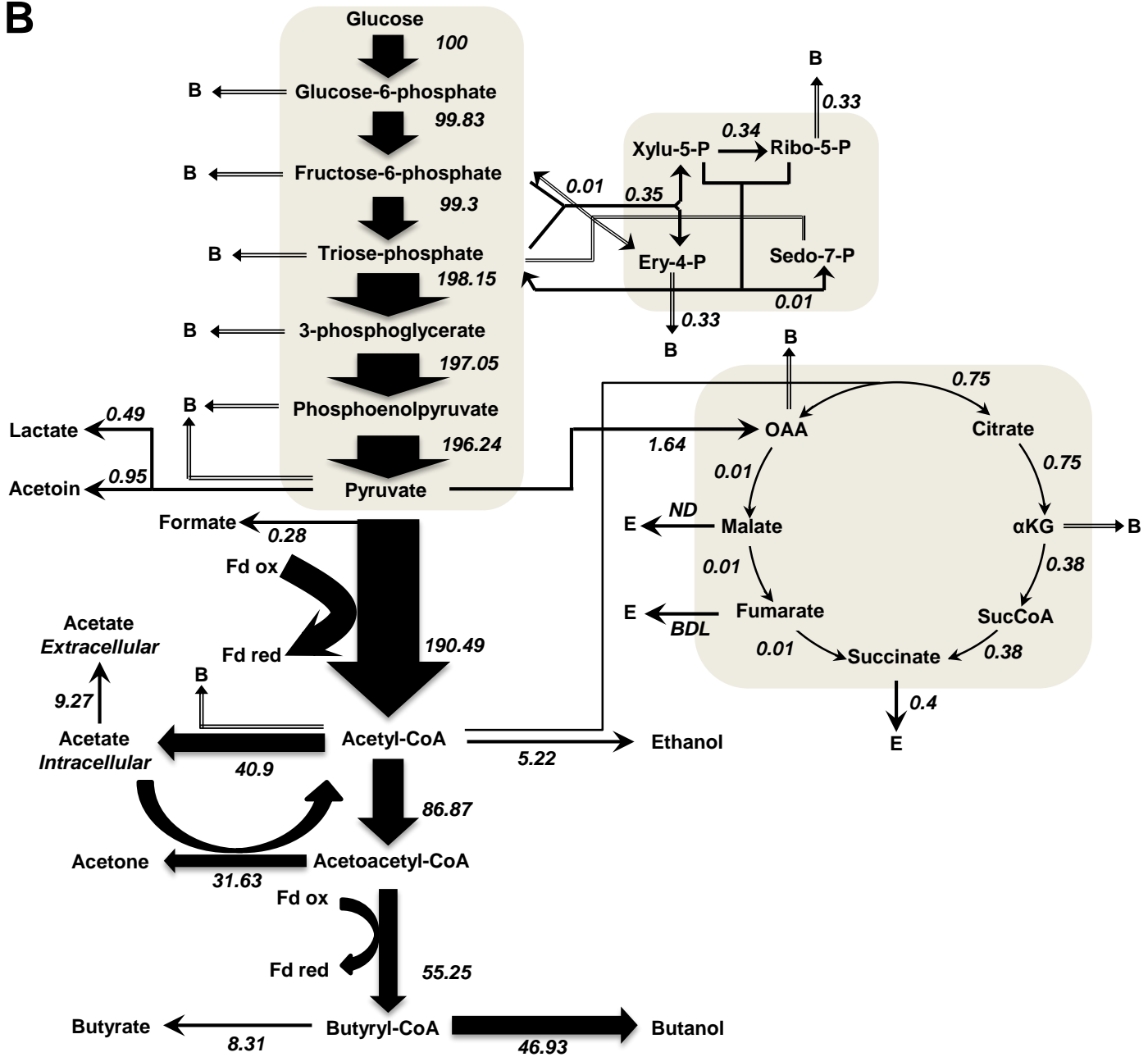


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C

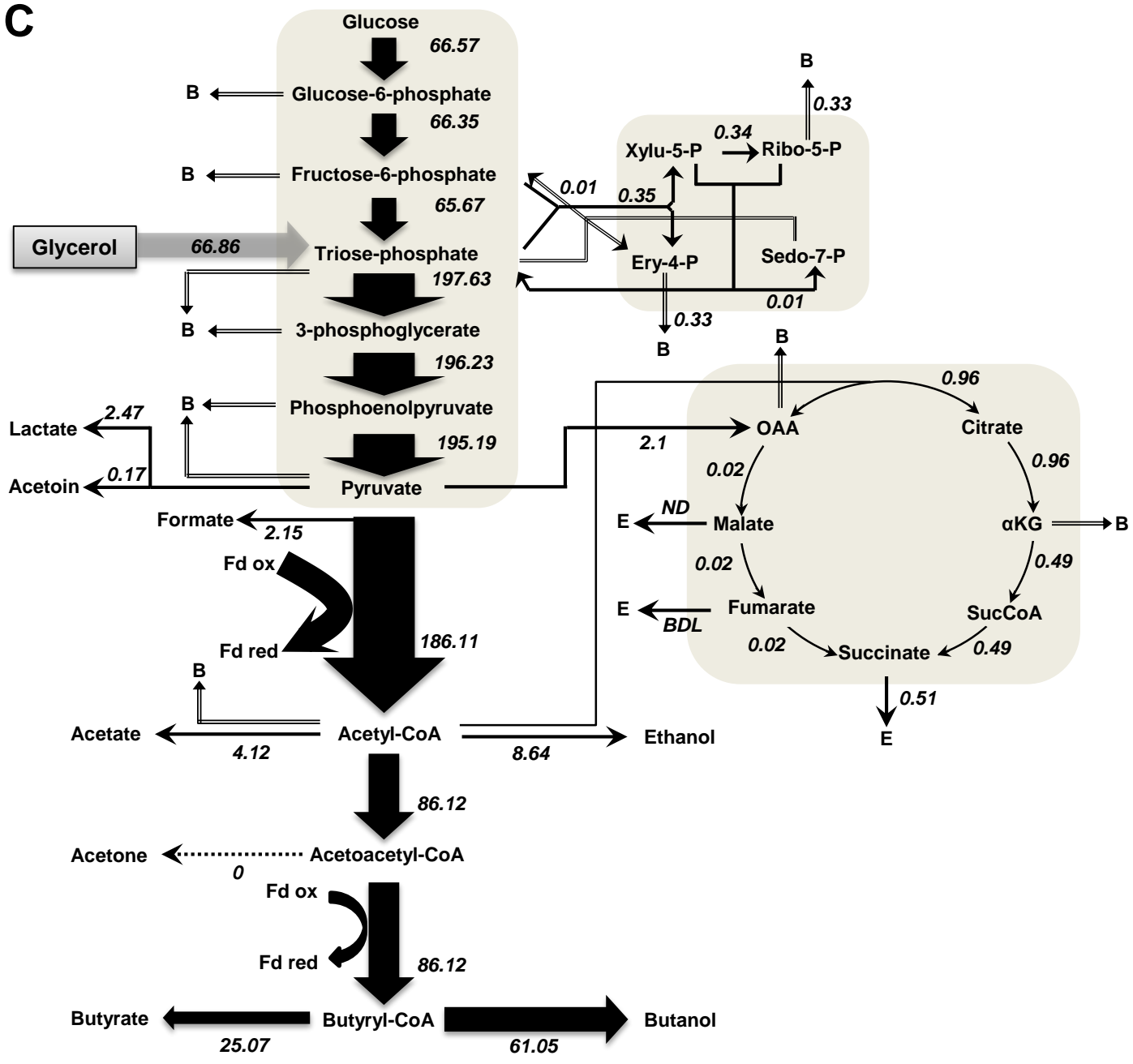


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