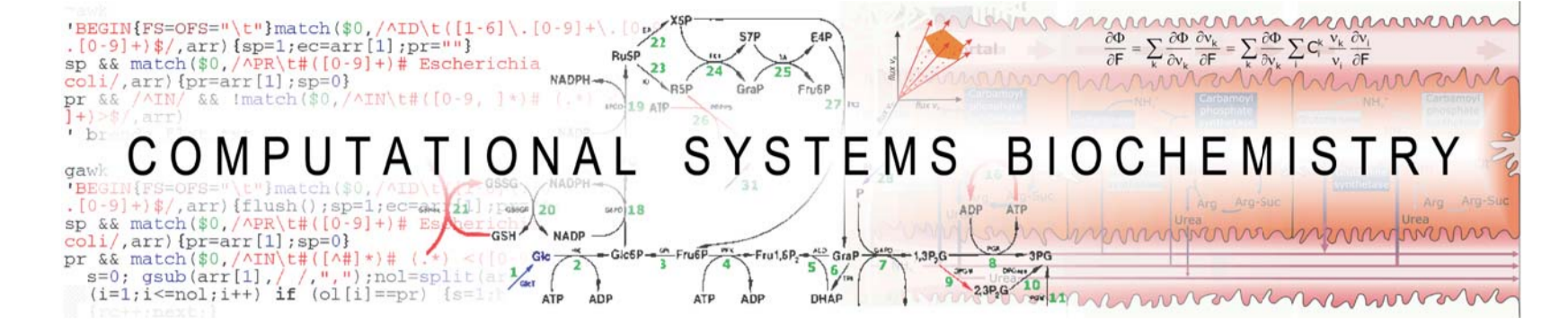


Are enzyme costs minimized in evolution?

Enzyme size, efficiency and turnover as a mean for better flux predictions in FBA

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Summary. Presumably, the costs of enzymes are minimized in the course of evolution, e.g. enzymes are as small (to minimize the effort for protein synthesis), as efficient (to minimize the necessary amount of enzymes for the required catalyzed flux), and with as low a degradation rate (to minimize the need to replenish the enzyme) as possible. However, as other factors (e.g. effective regulation

mechanisms) are critical for the cell's fitness it remains to be proved that costs have indeed an impact on the prevalence of enzymes throughout evolution. Here, the sequence lengths of the enzymes for central carbohydrate pathways is compared for selected classes in the tree of life in order to investigate the evolutionary preference of small enzymes. To assess the role of the different

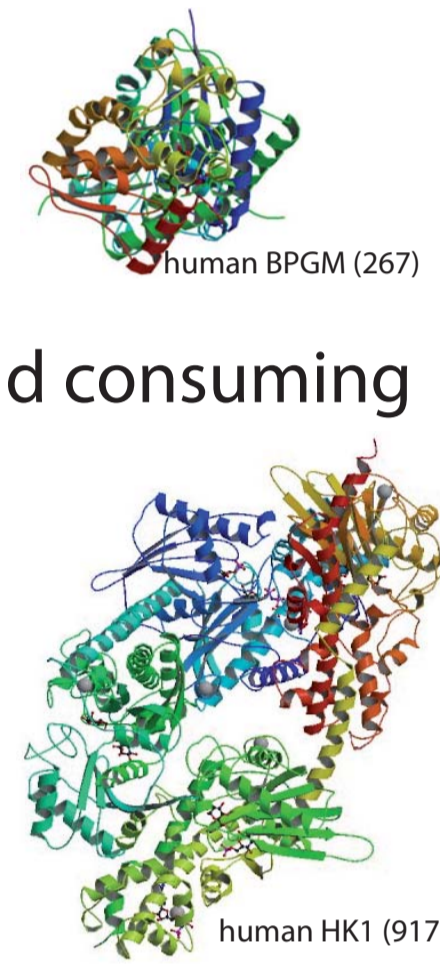
cost components the effect on the cost estimates is analyzed for two alternative pathways which yield glyceraldehyde phosphate and energy equivalents from activated glucose. As a concluding prospect, a flux-balance optimization procedure based on the minimization of required enzyme costs is presented.

Sequence length (by active site)

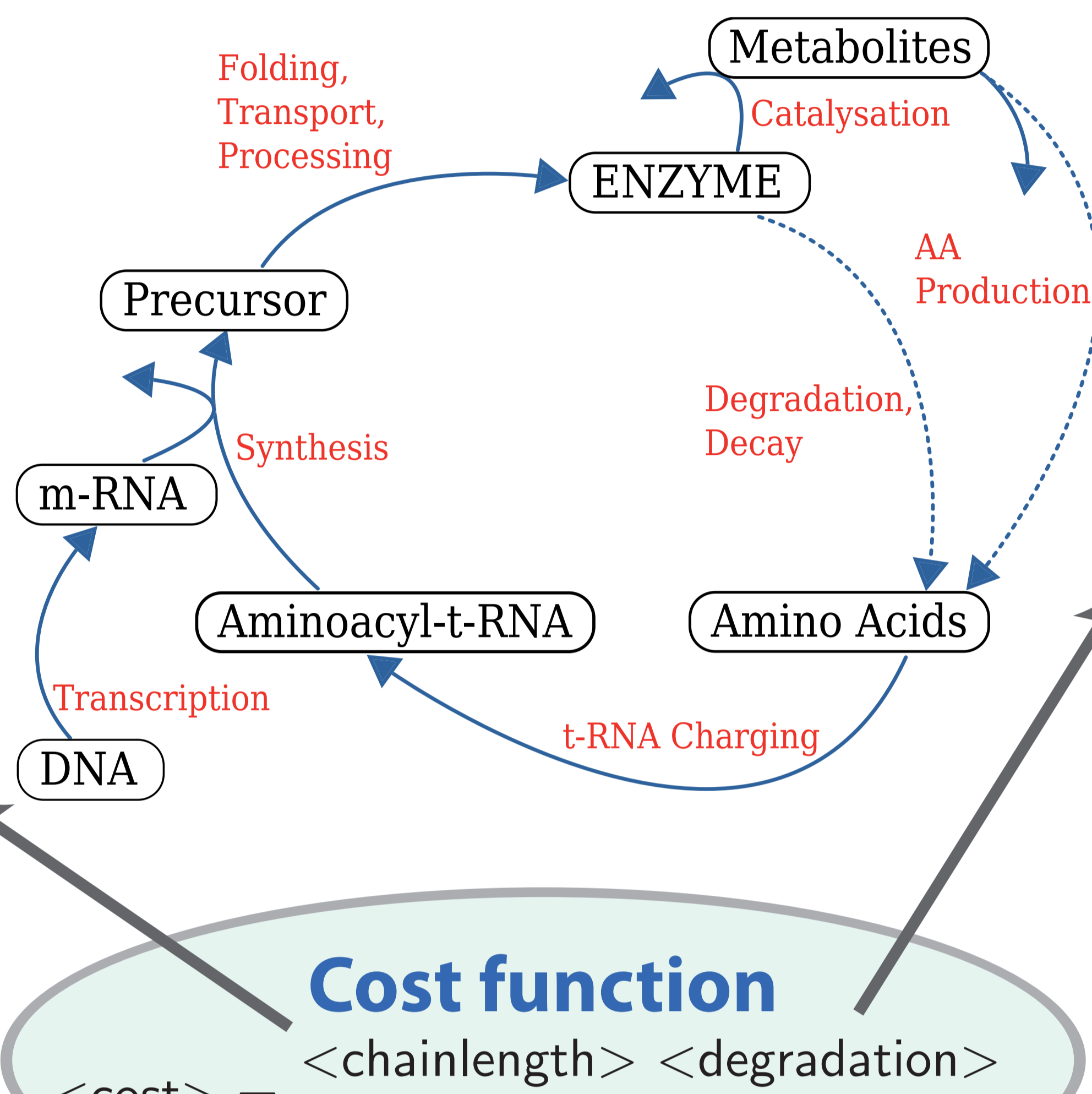
- fits perfectly in cost concept: **Investment**
- protein synthesis is major cell's energy consumer
- evolutionary pressure on protein size

Constraints:

- Stable folding
- Protective sequences (Protrypsin)
- Coupling of free energy producing and consuming chemical reaction
- Membrane position
- Allosteric mechanisms
- Stable binding of prosthetic groups
- Regulatory sites
- Reaction chamber (proteasome)



Data availability: perfect (uniprot+PDB)



Degradation rate

fits perfectly in cost concept: **Depreciation**

Constraints:

- Degradation very important regulation mechanism
- Protein modification/aging inevitable (radicals)
- Elimination of faulty enzymes (Ubiquitin)

Data availability: very poor, only few studies (e.g. [4])



Efficiency (turnover number)

fits relatively well in cost concept: **Productivity**

Constraints:

- Diffusion time
- Reaction mechanism

Problems:

- Depends on conditions
- Enzyme concentration and turnover not linear

Data availability: good (BRENDA) but large deviations

Cost function

$$\langle \text{cost} \rangle = \frac{\langle \text{chainlength} \rangle \langle \text{degradation} \rangle}{\langle \text{efficiency} \rangle \langle \text{specificity} \rangle}$$

Specificity (relative)

fits less well in cost concept: **Intensity of Labour**

- turnover of alternative reactants reduces efficiency
- Binding of chemically similar substances inevitable
- Essential side functions (e.g. stowaways)

Problem:

- depends on concentrations of alternative substrates

Typical: tradeoff between specificity and efficiency

Data availability: mediocre (BRENDA)



Comparison of chain lengths

- Selected pathways, all species, data from KEGG [6].
- Hierarchic arithmetic average chain lengths with respect to KEGG 4-level taxonomy used.
- Percentage (+/-) of average chain lengths of first category with respect to second shown.
- Results support: *intensive evolutionary development correlates with shorter chains.*

Pathway	Glycolysis										TCA										Pent-P ox										Pent-P non-ox									
	Reaction	Enzyme	Chain Length					
Eukaryotes vs. Prokaryotes	14	31	82	15	9	8	3	3	17	6	5	11	2	57	14	11	54	2	12	70	-3	6	44	-11	6	-3	33	-18	51	39	6	9	16							
Animals vs. Plants	-6	-15	53	-13	-1	-11	-4	-18	-15	-5	-4	-4	-1	-4	-6	7	6	2	-7	-3	-12	54	-15	-9	-14	-9	-11	-20	8	-8	-16									
Animals vs. Fungi	-2	-2	-5	-2	12	-1	3	-6	5	7	7	7	-5	-5	1	-15	-4	-2	-3	-2	6	22	-2	5	-1	1	-2	-1	4	22	-5									
Animals vs. Protists	9	-15	18	-12	-1	4	2	-12	12	-8	4	8	-5	1	-17	8	-2	1	-4	-3	-8	2	-15	-21	-37	-4	-9	-1	-3	25	-7									
Plants vs. Fungi	5	15	37	12	10	11	7	15	22	11	12	11	-3	-10	6	7	-10	0	5	2	20	-21	15	15	11	10	24	4	33	13										
Plants vs. Protists	16	0	-23	1	-2	8	6	8	32	-3	8	12	4	-5	-12	1	-8	3	3	1	5	-34	0	-13	27	5	3	23	5	36	10									
Fungi vs. Protists	10	-13	23	-10	-11	-3	-1	-6	8	-13	-3	1	-1	-5	-17	-6	2	3	-2	-1	-13	-17	-13	25	37	-5	-7	-1	2	-2										
Vertebrates vs. Arthropods	31	-1	-4	-1	1	1	-1	1	35	36	0	-2	-11	6	1	4	-3	8	5	-1	-2	25	-1	0	-13	2	4	19	-1	31	-5									
Vertebrates vs. Echinoderms	48	2	-5	-3	1	1	-1	-1	-40	-24	19	-2	-27	-1	-27	-1	-1	12	0	9	-24	2	35	-14	4	7	-1	-1	1	-6										
Vertebrates vs. Lancelets	39	-2	5	14	-2	-1	1	-1	-51	15	66	-5	35	39	-4	-3	1	-29	-5	2	1	-2	-2	-16	3	4	-5	0	11	4										
Mammals vs. Birds	-7	-6	1	-5	7	-3	-14	4	2	0	-1	-1	0	3	-3	-21	-1	-3	1	-6	-27	-6	8	10	10	-10	17	-2	5	7										
Mammals vs. Amphibians	-8	-6	-1	-5	6	-3	6	4	2	0	2	-2	49	1	-2	7	-20	-2	-3	-9	4	-6	1	10	0	-2	-2	30	14											
Mammals vs. Fishes	-10	-6	1	-4	6	-3	-1	4	2	6	1	-2	0	3	0	-1	-5	3	-2	3	-6	4	-6	-2	138	4	0	-2	-2	11	18									
Bacteria vs. Archaea	41	-2	27	4	14	-1	-1	-10	3	-1	1	43	26	4	1	-15	3	52	16	0	21	87	34	76	77	145	3	1	11											
Protists vs. Prokaryotes	5	41	74	-11	14	7	-1	5	-6	11	5	-3	61	14	25	49	4	10	71	-3	5	64	-4	25	30	34	55	42	5	7	16									
Protists vs. Bacteria	5	20	76	-11	11	1	0	6	-1	10	5	-3	60	11	23	48	5	8	71	-10	5	61	-9	25	30	17	22	43	4	6	10									
Protists vs. Archaea	70	72	13	16	15	-1	-5	-11	13	4	-2	129	40	28	49	-10	11	160	5	4	95	70	57	115	40	7	7	22												

Cost comparison Glycolysis vs. Pentose phosphate pathway

- Glycolysis versus Pentose-phosphate
- Substrate: glucose-6-phosphate; product: glyceraldehyde phosphate
- Data drawn from KEGG [6], BRENDA [5], and [4].
- Four Cost estimates: flux only, plus chain lengths considered, plus turnover numbers considered, plus degradation considered
- Glycolysis cheaper — main factor: turnover numbers

Reaction	flux	chain length	turnover number	degrad. rate	Enzyme cost for the reaction is expensive (red) / cheap (green)		
					cf/n	d	dcf/n
Glycolysis: 1 Glc6P + 1 ATP/ADP → 2 GraP							
GPI Glc6P → Fru6P	1	558	558	2456	0.23	0.17	0.04
PFK Fru6P → Fru1,6P	1	780	780	388	2.01	2.65	5.33
FBA Fru1,6P → GraP + DHAP	1	364	364	393.1	0.93	1.62	1.5
TPI GraP → DHAP	1	249	249	7512.5	0.03	0.17	0.01
∑	4		1951		3.2		6.9
Pentose-phosphate: 1 Glc6P + 0.67 ATP/ADP → 1.67 GraP + 1 CO₂ + 2 NADPH/NADP							
GPI Glc6P → bGlu6P	1	558	558	2456	0.23	0.17	0.04
G6PD bGlu6P → Glc1,5LacP	1	515	515	428.1	1.2	1.09	1.32
PGLS Glc1,5LacP → 6PG	1	258	258	(809)	0.32	(1)	0.32
PGD 6PG → Ru5P	1	483	483	15.9	30.43	0.24	7.33
PFK Fru6P → Fru1,6P	0.67	780	520	388	1.34	2.65	3.56
FBA Fru1,6P → GraP + DHAP	0.67	364	242.7	393.1	0.62	1.62	1
TPI GraP → DHAP	0.67	249	166	7512.5	0.02	0.17	0.004
RPE Ru5P → X5P	0.67	228	152	2609.6	0.06	(1)	0.06
rpiA Ru5P → R5P	0.33	311	103.7	25	4.15	(1)	4.15
tkt1 X5P + R5P → GraP + S7P	0.33	626	208.7	23.5	8.89	(1)	8.89
tal GraP + S7P → Fru6P + E4P	0.33	337	112.3	8.1	13.83	(1)	13.83
tkt2 X5P + E4P → Fru6P + GraP	0.33	596	198.7	34.5	5.76	(1)	5.76
∑	8		3518		66.8		46.2

Using cost as an objective for FBA

- Adaptation of the flux minimization principle [3].
- Objective: absolute flux value multiplied by specific enzyme cost.
- Linear optimization problem., implemented in [2], see [1] for details.

Minimize $\sum_{i=1}^n c_i v_i$

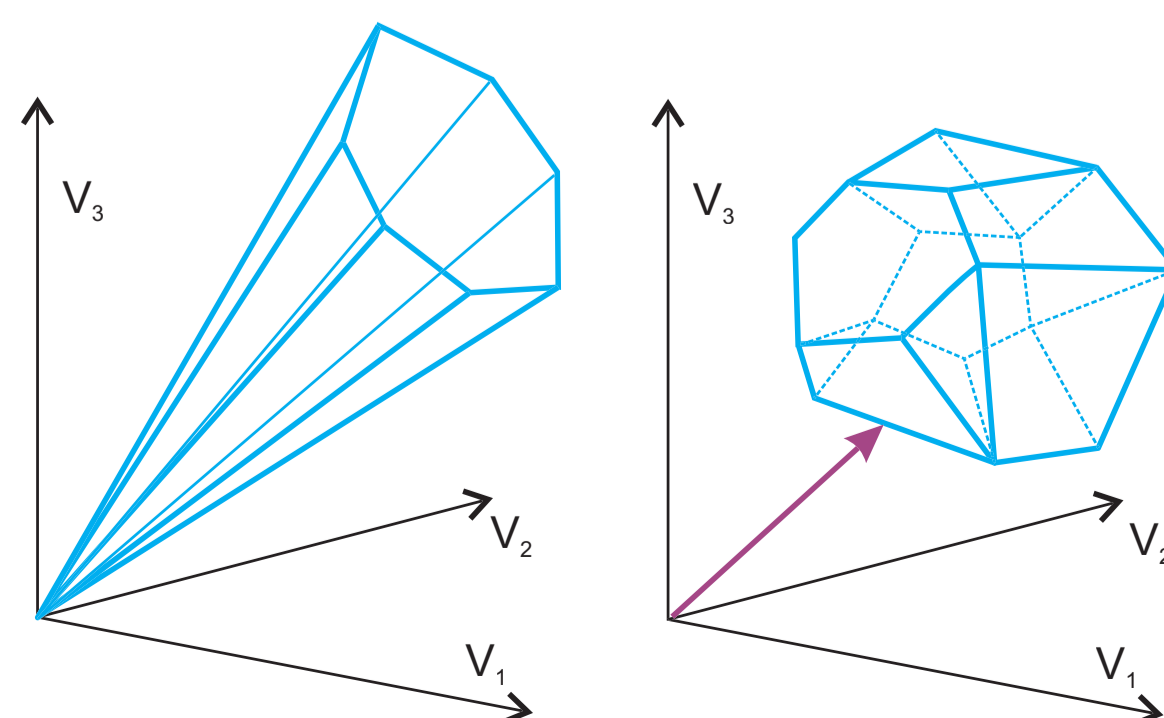
Subject to $\sum_{i=1}^n s_{ij} v_j = 0, \quad \forall j \text{ internal}$

$v_i = L_i, \quad \forall i \text{ target flux}$

(c_1, \dots, c_n) enzyme costs vector

(v_1, \dots, v_n) flux vector

$(s_{ij})_{ij}$ stoichiometric matrix



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