Supplementary Material for: Temporal microbiome road-maps guided by perturbations

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Figure S1: Human gut microbiome sample distribution

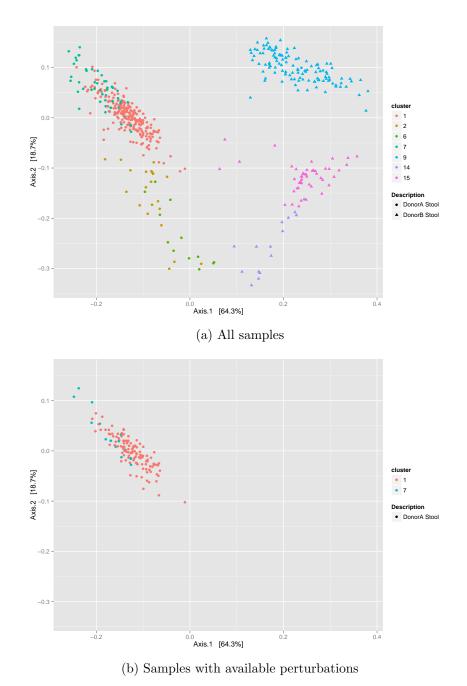


Figure S1: Sample distribution in clusters from human gut microbiome dataset. Before (a) and after (b) removing samples without perturbations.

Figure S2: Vaginal flora microbiome sample distribution

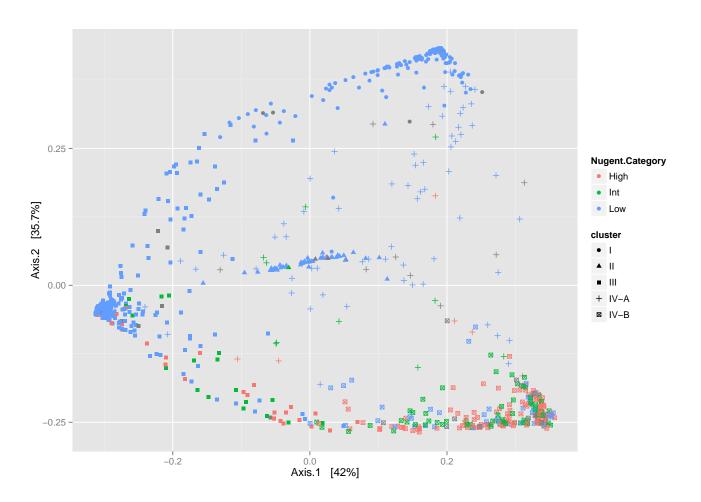


Figure S2: Sample distribution in clusters from vaginal flora microbiome dataset. Distribution of Nugent score per sample, being 'High' value which indicates the greatest risk for the Bacterial Vaginosis disease, concentrated mainly in cluster IV-B.

Table S1: transition probabilities in Vaginal flora microbiome dataset

Table S1: Global relative transition probabilities in Vaginal flora microbiome dataset. Input state in rows, output state in columns. Each row sums 1.

States	I	II	III	IV-A	IV-B
I	0.8418	0.0000	0.0759	0.0696	0.0127
II	0.0000	0.8929	0.0357	0.0714	0.0000
III	0.0273	0.0055	0.8579	0.0437	0.0656
IV-A	0.1364	0.0341	0.1932	0.5568	0.0795
IV-B	0.0127	0.0000	0.0928	0.0169	0.8776

Figure S3: Transition diagram vaginal flora microbiom

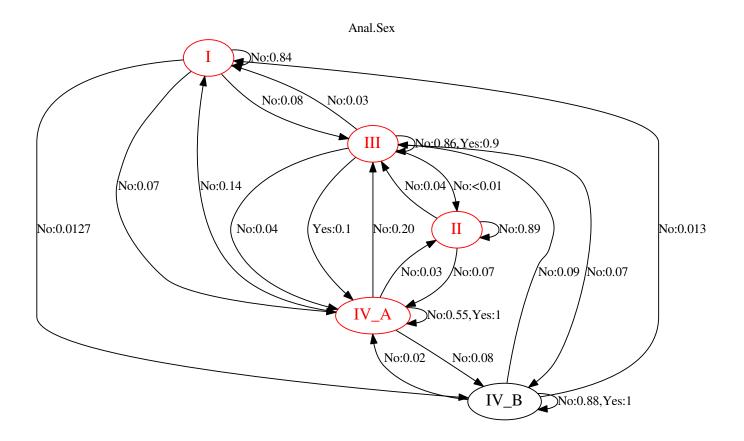


Figure S3: Transition diagram of vaginal flora microbiome with anal sex as perturbation. Goal (healthy) states highlighted in red.

Table S2: Combined MDP policy in Human Gut

Table S2: Quantity of nutritional intakes to reach *clus* 7 as the desired microbiome state. Rows represent external perturbations that could influence the human gut microbiome. Columns represent the microbiome states. In each cell, the policy (low/medium/high) from all Markov systems, one per external perturbation.

Perturbation \ State	clus1	clus 7
Calcium	Medium	Low
Calorie	Low	Medium
Carbohydrates	Medium	Low
Cholesterol	Low	Medium
Fat	Low	Medium
Fiber	High	Low
Protein	Low	Medium
Saturated fat	Low	Medium
Sodium	Low	Medium
Sugar	High	High