

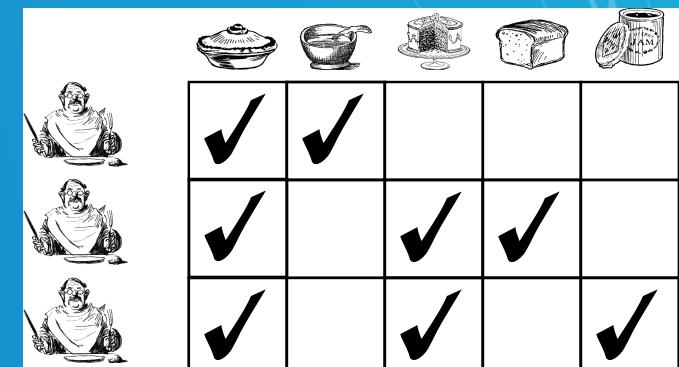
Probabilistic Graphical Models

Bayesian Nonparametrics: Indian Buffet Process

Eric Xing

Lecture 24, April 15, 2020

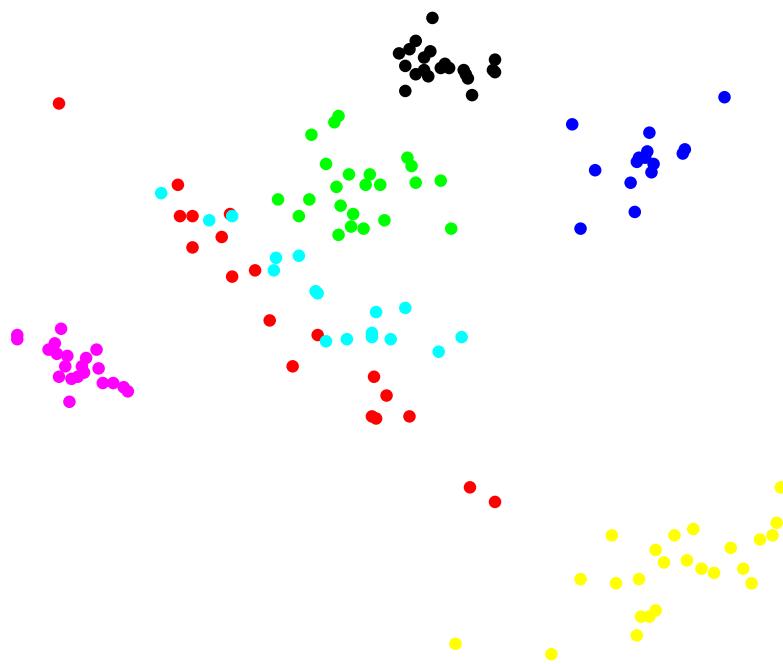
Reading: see class homepage





Recap of last lecture

- ❑ Dirichlet process: a distribution over discrete probability distributions with infinitely many atoms.
- ❑ Can be used to create a *nonparametric* version of a *finite mixture model*.





Recap of last lecture

- We can think of the Dirichlet process in a number of ways:
 - The **infinite limit** of a Dirichlet distribution.
 - A rich-gets-richer predictive distribution over the next data point (**Chinese restaurant process, Polya urn scheme**).
 - An iterative procedure for generating samples from the Dirichlet process – the **stick breaking representation**.





Limitations of a simple mixture model

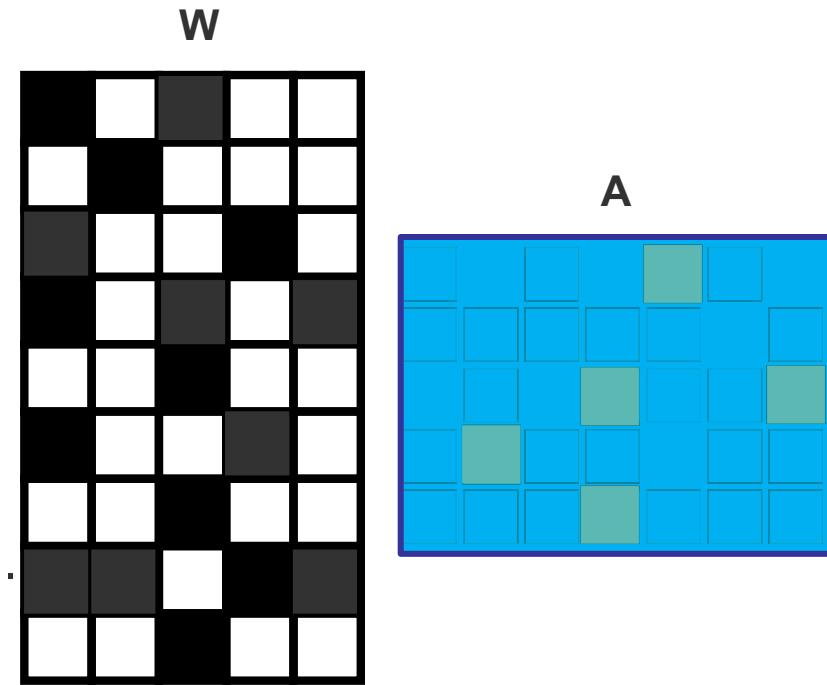
- The Dirichlet distribution and the Dirichlet process are great if we want to cluster data into non-overlapping clusters.
- However, DP/Dirichlet mixture models cannot share features (i.e., cluster centroids, prototypes) between clusters.
- In many applications, data points exhibit properties of multiple latent features
 - Images contain multiple objects.
 - Actors in social networks belong to multiple social groups.
 - Movies contain aspects of multiple genres.





Latent variable models

- Latent variable models allow each data point to exhibit *multiple latent features*, to *varying degrees*.
- Example: Factor analysis
 - $\mathbf{X} = \mathbf{WA}^T + \varepsilon$
 - Rows of \mathbf{A} = latent features
 - Rows of \mathbf{W} = data-point-specific weights for these features
 - ε = Gaussian noise.
- Example: LDA
 - Each document represented by a *mixture* of features.



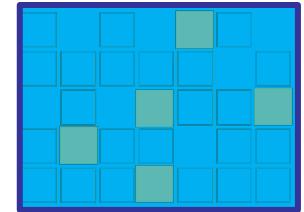
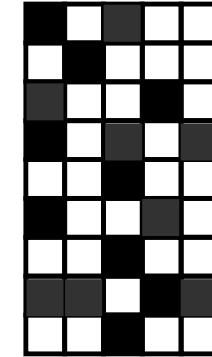


Infinite latent feature models

- Problem: How to choose the number of features?
- Example: Factor analysis

$$\mathbf{X} = \mathbf{W}\mathbf{A}^T + \varepsilon$$

- Each column of \mathbf{W} (and row of \mathbf{A}) corresponds to a feature.
- Question: Can we make the number of features *unbounded* a posteriori, as we did with the DP?
- Solution: allow *infinitely many* features a priori – i.e. let \mathbf{W} (or \mathbf{A}) have infinitely many columns (rows).
- Problem: We can't represent infinitely many features!
- Solution: make our infinitely large matrix *sparse*.



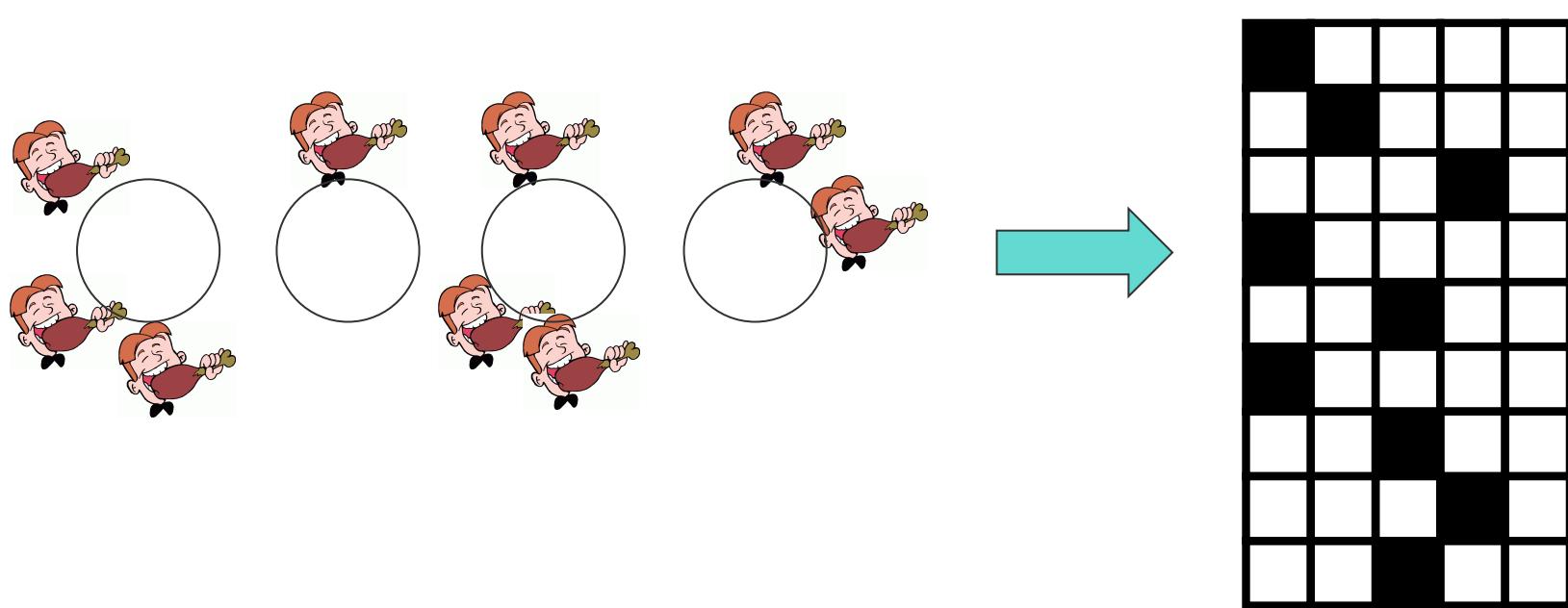
Griffiths and Ghahramani, 2006





Recall the CRP: a distribution over indicator matrices

- Recall that the CRP gives us a distribution over *partitions* of our data.
- We can represent this as a distribution over *binary (**indicator**) matrices*, where each row (which is a “one-hot vector”) corresponds to a data point, and each column to a cluster.





A sparse, finite latent variable model

- We want a *sparse* model – so let

$$\mathbf{X} = \mathbf{W}\mathbf{A}^T + \epsilon$$

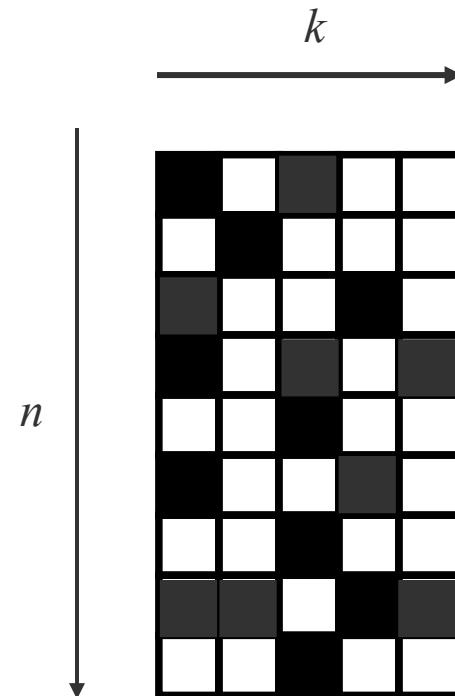
$$\mathbf{W} = \mathbf{Z} \odot \mathbf{V}$$

for some sparse matrix \mathbf{Z} .

- Place a *beta-Bernoulli prior* on \mathbf{Z} :

$$\pi_k \sim \text{Beta}\left(\frac{\alpha}{K}, 1\right), k = 1, \dots, K$$

$$z_{nk} \sim \text{Bernoulli}(\pi_k), n = 1, \dots, N.$$





A sparse, finite latent variable model

- If we integrate out the π_k , the marginal probability of a matrix \mathbf{Z} is:

$$p(\mathbf{Z}) = \prod_{k=1}^K \int \left(\prod_{n=1}^N p(z_{nk} | \pi_k) \right) p(\pi_k) d\pi_k$$

$$= \prod_{k=1}^K \frac{B(m_k + \alpha/K, N - m_k + 1)}{B(\alpha/K, 1)}$$

$$= \prod_{k=1}^K \frac{\alpha}{K} \frac{\Gamma(m_k + \alpha/K)\Gamma(N - m_k + 1)}{\Gamma(N + 1 + \alpha/K)}$$

where

$$m_k = \sum_{n=1}^N z_{nk}$$

- This is *exchangeable* (doesn't depend on the order of the rows or columns)





A sparse, finite latent variable model

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where

$$m_k = \sum_{n=1}^N z_{nk}$$

- How is this sparse?

$$\begin{aligned} & m_k! (N - m_k)! / (N + 1)! \\ &= 1 * 2 * \dots * (N - m_k) / (m_k + 1) * (m_k + 2) * \dots * (N + 1) \\ &= (1 / (m_k + 1)) * (2 / (m_k + 2)) * \dots * ((N - m_k) / (N + 1)) \end{aligned}$$





An equivalence class of matrices

- We can naively take the infinite limit by taking K to infinity
- Because all the columns are equal in expectation, as K grows we are going to have more and more empty columns.
- We do not want to have to represent infinitely many empty columns!
- Define an *equivalence class* $[\mathbf{Z}]$ of matrices where the non-zero columns are all to the left of the empty columns.
- Let $l\text{o}f(\cdot)$ be a function that maps binary matrices to *left-ordered* binary matrices – matrices ordered by the binary number made by their rows.





Left-ordered matrices

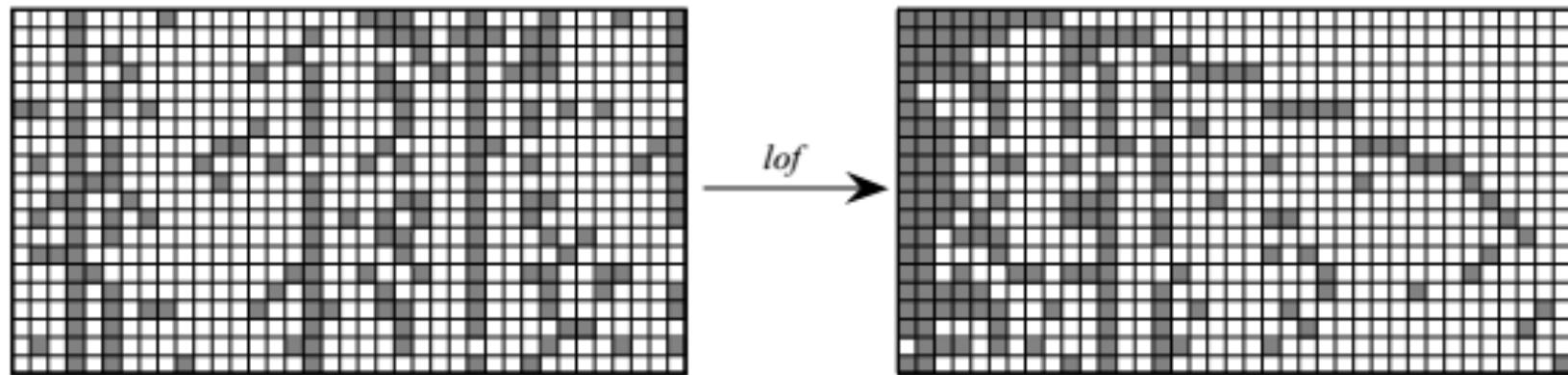


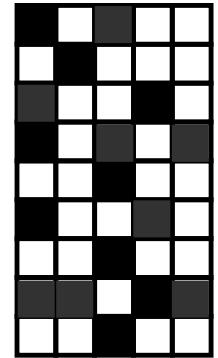
Figure 5: Binary matrices and the left-ordered form. The binary matrix on the left is transformed into the left-ordered binary matrix on the right by the function $lof(\cdot)$. This left-ordered matrix was generated from the exchangeable Indian buffet process with $\alpha = 10$. Empty columns are omitted from both matrices.

Image from Griffiths and Ghahramani, 2011





How big is the equivalence set?



- All matrices in the equivalence set $[\mathbf{Z}]$ are equiprobable (by exchangeability of the columns), so if we know the size of the equivalence set, we know its probability.
- Call the vector $(z_{1k}, z_{2,k}, \dots, z_{(n-1)k})$ the *history* of feature k at data point n (a number represented in binary form).
- Let K_h be the number of features possessing history h , and let K_+ be the total number of features with non-zero history.
- The total number of lof-equivalent matrices in $[\mathbf{Z}]$ is

$$\binom{K}{K_0 \cdots K_{2^N-1}} = \frac{K!}{\prod_{n=0}^{2^N-1} K_n!}$$





Probability of an equivalence class of finite binary matrices.

- If we know the size of the equivalence class $[\mathbf{Z}]$, we can evaluate its probability:

$$\begin{aligned} p([\mathbf{Z}]) &= \sum_{\mathbf{Z} \in [\mathbf{Z}]} p(\mathbf{Z}) \\ &= \frac{K!}{\prod_{n=0}^{2^N-1} K_n!} \prod_{k=1}^K \frac{\alpha}{K} \frac{\Gamma(m_k + \alpha/K) \Gamma(N - m_k + 1)}{\Gamma(N + 1 + \alpha/K)} \\ &= \frac{\alpha^{K_+}}{\prod_{n=1}^{2^N-1} K_n!} \frac{K!}{K_0! K^{K_+}} \left(\frac{N!}{\prod_{j=1}^N j + \alpha/K} \right)^K \\ &\cdot \prod_{k=1}^{K_+} \frac{(N - m_k)! \prod_{j=1}^{m_k-1} (j + \alpha/K)}{N!} \end{aligned}$$





Taking the infinite limit

- We are now ready to take the limit of this finite model as K tends to infinity:

$$\frac{\alpha^{K_+}}{\prod_{n=1}^{2^N-1} K_n!} \frac{K!}{K_0! K^{K_+}} \left(\frac{N!}{\prod_{j=1}^N j + \frac{\alpha}{K}} \right)^K \prod_{k=1}^{K_+} \frac{(N - m_k)! \prod_{j=1}^{m_k-1} (j + \frac{\alpha}{K})}{N!}$$

$\downarrow K \rightarrow \infty$

$$\frac{\alpha^{K_+}}{\prod_{n=1}^{2^N-1} K_n!}$$

1

$$\exp\{-\alpha H_N\}$$

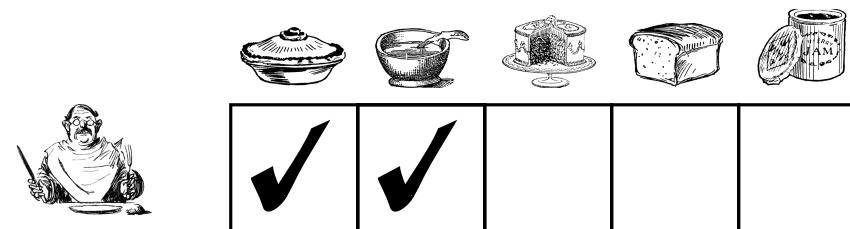
$$\prod_{k=1}^{K_+} \frac{(N - m_k)!(m_k - 1)!}{N!}$$





Predictive distribution: The Indian buffet process

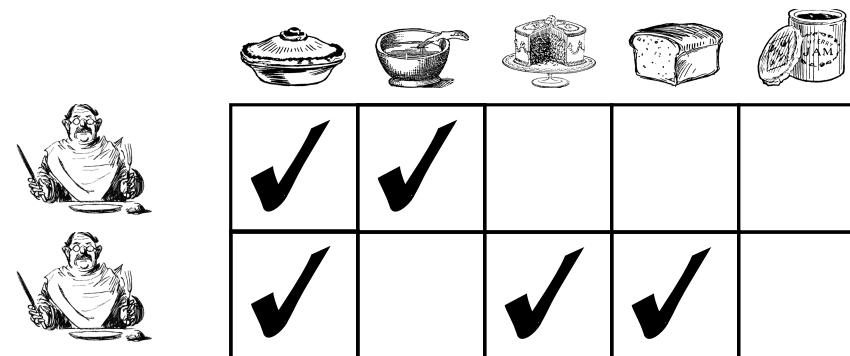
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 - A customer enters a restaurant with an infinitely large buffet
 - He helps himself to Poisson(a) dishes.





Predictive distribution: The Indian buffet process

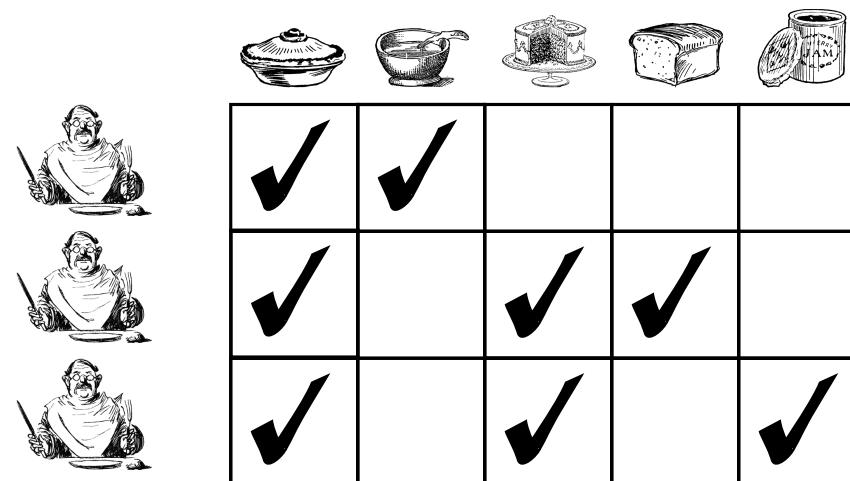
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 - The n^{th} customer enters the restaurant
 - He helps himself to each previously chosen dish with probability m_k / n
 - He then tries Poisson(a/n) new dishes





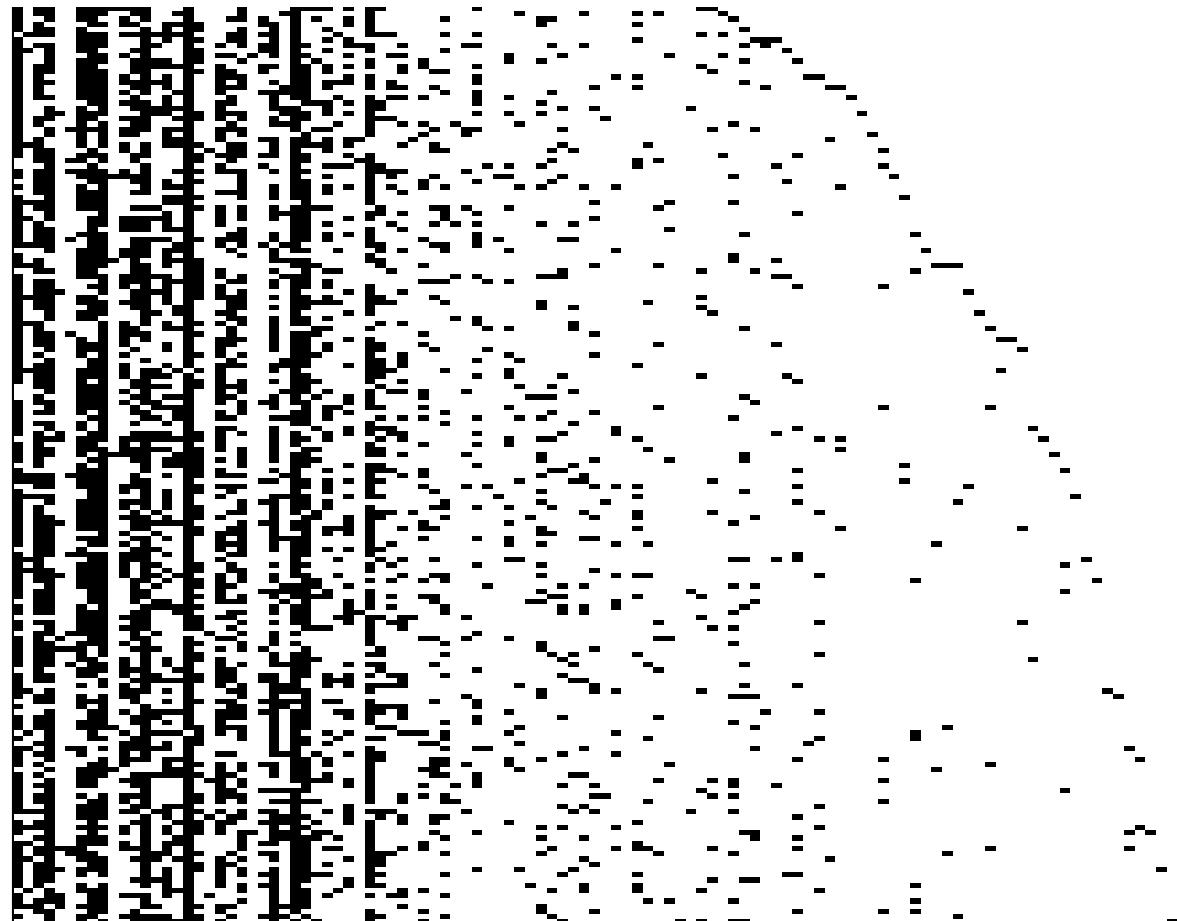
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Example





Proof that the IBP is lof-equivalent to the infinite beta-Bernoulli model

- What is the probability of a matrix \mathbf{Z} ?
- Let $K_1^{(n)}$ be the number of new features in the n^{th} row.

$$\begin{aligned} p(\mathbf{Z}) &= \prod_{n=1}^N p(\mathbf{z}_n | \mathbf{z}_{1:(n-1)}) \\ &= \prod_{n=1}^N \text{Poisson}\left(K_1^{(n)} \middle| \frac{\alpha}{n}\right) \prod_{k=1}^{K_+} \left(\frac{\sum_{i=1}^{n-1} z_{ik}}{n}\right)^{z_{nk}} \left(\frac{n - \sum_{i=1}^{n-1} z_{ik}}{n}\right)^{1-z_{nk}} \\ &= \prod_{n=1}^N \left(\frac{\alpha}{n}\right)^{K_1^{(n)}} \frac{1}{K_1^{(n)}!} e^{-\alpha/n} \prod_{k=1}^{K_+} \left(\frac{\sum_{i=1}^{n-1} z_{ik}}{n}\right)^{z_{nk}} \left(\frac{n - \sum_{i=1}^{n-1} z_{ik}}{n}\right)^{1-z_{nk}} \\ &= \frac{\alpha^{K_+}}{\prod_{n=1}^N K_1^{(n)}!} \exp\{-\alpha H_N\} \prod_{k=1}^{K_+} \frac{(N-m_k)!(m_k-1)!}{N!} \end{aligned}$$

- If we include the cardinality of $[\mathbf{Z}]$, this is the same as before





Properties of the IBP

- “Rich get richer” property – “popular” dishes become more popular.
- The number of nonzero entries for each row is distributed according to $\text{Poisson}(a)$ – due to exchangeability.
- Recall that if $x_1 \sim \text{Poisson}(a_1)$ and $x_2 \sim \text{Poisson}(a_2)$, then $(x_1 + x_2) \sim \text{Poisson}(a_1 + a_2)$
 - The number of nonzero entries for the whole matrix is distributed according to $\text{Poisson}(Na)$.
 - The number of non-empty columns is distributed according to $\text{Poisson}(aH_N)$





Building latent feature models using the IBP

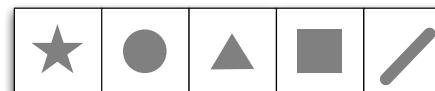
- We can use the IBP to build latent feature models with an unbounded number of features.
- Let each column of the IBP correspond to one of an *infinite* number of features.
- Each row of the IBP selects a *finite subset* of these features.
- The **rich-get-richer** property of the IBP ensures features are shared between data points.
- We must pick a *likelihood model* that determines **what the features look like** and **how they are combined**.





A linear Gaussian model

- General form of latent factor model: $\mathbf{X} = \mathbf{W}\mathbf{A}^T + \boldsymbol{\varepsilon}$
- Simplest way to make an infinite factor model:
 - Sample $\mathbf{W} \sim \text{IBP}(a)$
 - Sample $\mathbf{a}_k \sim \mathcal{N}(\mathbf{0}, \sigma_a^2 \mathbf{I})$
 - Sample $\boldsymbol{\varepsilon}_{nk} \sim \mathcal{N}(0, \sigma_\varepsilon^2)$





Infinite factor analysis

- Problem with linear Gaussian model: Features are “all or nothing” due to the binary “loading matrix” \mathbf{W} .
- Factor analysis: $\mathbf{X} = \mathbf{WA}^T + \varepsilon$
 - Rows of \mathbf{A} = latent features (Gaussian)
 - Rows of \mathbf{W} = data-point-specific **weights** for these features (Gaussian)
 - ε = Gaussian noise.
- Write
$$\mathbf{W} = \mathbf{Z} \odot \mathbf{V}$$
 - $\mathbf{Z} \sim \text{IBP}(\alpha)$
 - $\mathbf{V} \sim \mathcal{N}(0, \sigma_v^2)$
 - $\mathbf{A} \sim \mathcal{N}(0, \sigma_A^2)$





A binary model for latent networks

- Motivation: Discovering latent causes for observed binary data
- Example:
 - Data points = patients
 - Observed features = presence/absence of symptoms
 - Goal: Identify biologically plausible “latent causes” – e.g. illnesses.
- Idea:
 - Each latent feature is associated with a set of symptoms
 - The more features a patient has that are associated with a given symptom, the more likely that patient is to exhibit the symptom.

Wood et al, 2006





A binary model for latent networks

- We can represent this in terms of a *Noisy-OR* model:

$$\mathbf{Z} \sim \text{IBP}(\alpha)$$

$$y_{dk} \sim \text{Bernoulli}(p) \quad \begin{matrix} d^{\text{th}} \text{ observed symptom,} \\ k^{\text{th}} \text{ latent disease} \end{matrix}$$

$$p(x_{nd} = 1 | \mathbf{Z}, \mathbf{Y}) = 1 - (1 - \lambda)^{\mathbf{z}_n} \mathbf{y}_d^T (1 - \epsilon)$$

- Intuition:
 - Each patient has a set of latent causes, as indicated by Z
 - Each latent cause (disease) k exhibit a symptom d with a Bernoulli rate
 - For each symptom,, we toss a coin with probability λ for each latent cause that is “on” for that patient and associated with that feature, plus an extra coin with probability ϵ .
 - If any of the coins land heads, we exhibit that feature.





Inference in the IBP

- ❑ Recall inference methods for the DP:
 - ❑ Gibbs sampler based on the exchangeable model.
 - ❑ Gibbs sampler based on the underlying Dirichlet distribution
 - ❑ Variational inference
 - ❑ Particle filter.
- ❑ We can construct analogous samplers for the IBP





Inference in the restaurant scheme

- Recall the exchangeability of the IBP means we can treat any data point as if it's our last.
- Let K_+ be the total number of used features, excluding the current data point.
- Let Θ be the set of parameters associated with the likelihood – eg the Gaussian matrix \mathbf{A} in the linear Gaussian model
- The prior probability of choosing one of these features is m_k / N
- The posterior probability is proportional to
$$p(z_{nk} = 1 | \mathbf{x}_n, \mathbf{Z}_{-nk}, \Theta) \propto m_k f(\mathbf{x}_n | z_{nk} = 1, \mathbf{Z}_{-nk}, \Theta)$$
$$p(z_{nk} = 0 | \mathbf{x}_n, \mathbf{Z}_{-nk}, \Theta) \propto (N - m_k) f(\mathbf{x}_n | z_{nk} = 0, \mathbf{Z}_{-nk}, \Theta)$$
- In some cases, we can integrate out Θ , otherwise we must sample this.





Inference in the restaurant scheme

- In addition, we must propose adding new features.
- Metropolis Hastings method:
 - Let K_{old}^* be the number of features appearing only in the current data point.
 - Propose $K_{new}^* \sim \text{Poisson}(a/N)$, and let \mathbf{Z}^* be the matrix with K_{new}^* features appearing only in the current data point.
 - With probability

$$\min \left(1, \frac{f(\mathbf{x}_n | \mathbf{Z}^*, \Theta)}{f(\mathbf{x}_n | \mathbf{Z}, \Theta)} \right)$$

accept the proposed matrix.





Beta processes and the IBP

- ❑ Recall the relationship between the Dirichlet process and the Chinese restaurant process:
 - ❑ The Dirichlet process is a prior on probability measures (distributions)
 - ❑ We can use this probability measure as cluster weights in a clustering model
 - cluster allocations are i.i.d. given this distribution.
 - ❑ If we integrate out the weights, we get an *exchangeable* distribution over partitions of the data – the **Chinese restaurant process**.
- ❑ De Finetti's theorem tells us that, if a distribution X_1, X_2, \dots is *exchangeable*, there **must** exist a measure conditioned on which X_1, X_2, \dots are i.i.d.





Beta processes and the IBP

- Recall the finite beta-Bernoulli model:

$$\pi_k \sim \text{Beta}\left(\frac{\alpha}{K}, 1\right)$$

$$z_{nk} \sim \text{Bernoulli}(\pi_k)$$

- The z_{nk} are i.i.d. given the π_k , but are exchangeable if we integrate out the π_k .
- The corresponding distribution for the IBP is the *infinite limit* of the beta random variables, as K tends to infinity.
- This distribution over discrete measures is called the **beta process**.
- Samples from the beta process have infinitely many atoms with masses between 0 and 1.





Posterior distribution of the beta process

- **Question:** Can we obtain the posterior distribution of the column probabilities in closed form?
- **Answer:** Yes!
 - Recall that each atom of the beta process is the infinitesimal limit of a $\text{Beta}(a/K, 1)$ random variable.
 - Our counts of observations for that atom are a $\text{Binomial}(\pi_k, N)$ random variable.
 - We know the beta distribution is conjugate to the Binomial, so the posterior is the infinitesimal limit of a $\text{Beta}(a/K+m_k, N+1-m_k)$ random variable.

Theorem: Let X_1, X_2, \dots, X_n be independent Bernoulli random variables, each with the same parameter p . Then the sum $X = X_1 + \dots + X_n$ is a binomial random variable with parameters n and p .





A stick-breaking construction for the beta process

- We can construct the beta process using the following stick-breaking construction:
- Begin with a stick of unit length.
- For $k=1,2,\dots$
 - Sample a $\text{Beta}(\alpha, 1)$ random variable μ_k .
 - Break off a fraction μ_k of the stick. This is the k^{th} atom size.
 - Throw away *what's left* of the stick.
 - Recurse on the part of the stick that you broke off

$$\pi_k = \prod_{j=1}^k \mu_j \quad \mu_j \sim \text{Beta}(\alpha, 1)$$

- Note that, unlike the DP stick breaking construction, the atoms will *not* sum to one.





Inference in the stick-breaking construction

- We can also perform inference using the stick-breaking representation
 - Sample $\mathbf{Z}|\boldsymbol{\pi}, \Theta$
 - Sample $\boldsymbol{\pi}|\mathbf{Z}$
- The posterior for atoms for which $m_k > 0$ is beta distributed.
- The atoms for which $m_k = 0$ can be sampled using the stick-breaking procedure.
- We can use a *slice sampler* to avoid representing all of the atoms, or using a fixed truncation level.

Teh et al, 2007





A two-parameter extension

- In the IBP, the parameter a governs both the *number of nonempty columns* and the *number of features per data point*.
- We might want to decouple these properties of our model.
- Reminder: We constructed the IBP as the limit of a finite beta-Bernoulli model where

$$\pi_k \sim \text{Beta}\left(\frac{\alpha}{K}, 1\right)$$

$$z_{nk} \sim \text{Bernoulli}(\pi_k)$$

- We can modify this to incorporate an extra parameter:

$$\pi_k \sim \text{Beta}\left(\frac{\alpha\beta}{K}, \beta\right)$$

$$z_{nk} \sim \text{Bernoulli}(\pi_k)$$

Sollich, 2005





A two-parameter extension

- Our restaurant scheme is now as follows:
 - A customer enters a restaurant with an infinitely large buffet
 - He helps himself to Poisson(a) dishes.
 - The n^{th} customer enters the restaurant
 - He helps himself to each dish with probability $m_k / (\beta+n-1)$
 - He then tries Poisson($a\beta / (\beta+n-1)$) new dishes
- Note
 - The number of features per data point is still marginally Poisson(a).
 - The number of non-empty columns is now

$$\text{Poisson}\left(\alpha \sum_{n=1}^N \frac{\beta}{\beta+n-1}\right)$$

- We recover the IBP when $\beta = 1$.



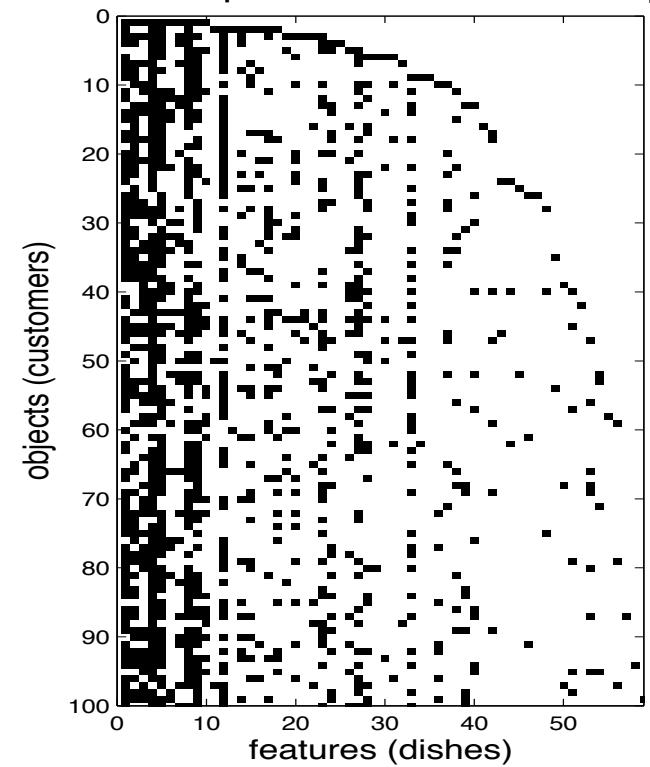


Two parameter IBP: examples

from IBP

2

Prior sample from IBP with $\alpha=10 \beta=1$



ies)

Prior sample from IBP with $\alpha=10 \beta=5$

objects (customers)

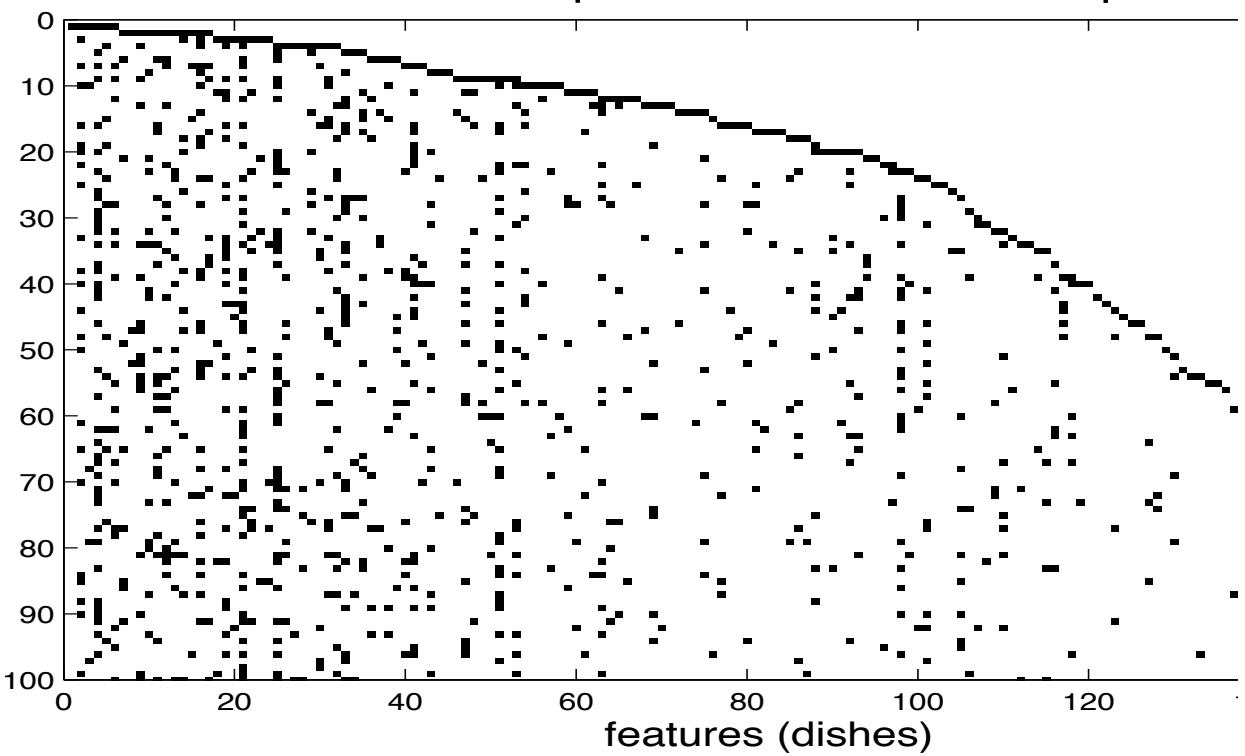


Image from Griffiths and Ghahramani, 2011





Other distributions over infinite, exchangeable matrices

- Recall the beta-Bernoulli process construction of the IBP.
- We start with a beta process – an infinite sequence of values between 0 and 1 that are distributed as the infinitesimal limit of the beta distribution.
- We combine this with a Bernoulli process, to get a binary matrix.
- If we integrate out the beta process, we get an exchangeable distribution over binary matrices.
- Integration is straightforward due to the beta-Bernoulli conjugacy.
- **Question:** Can we construct other infinite matrices in this way?





The infinite gamma-Poisson process

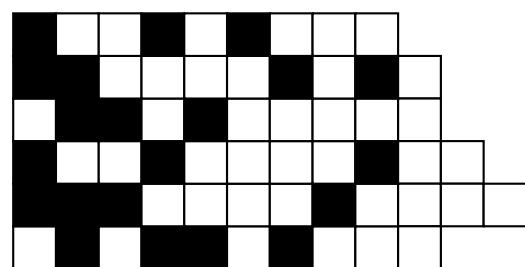
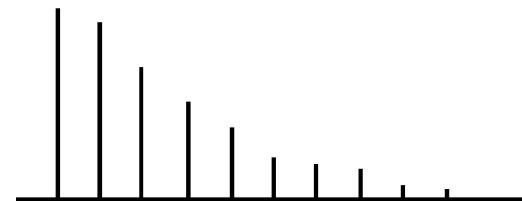
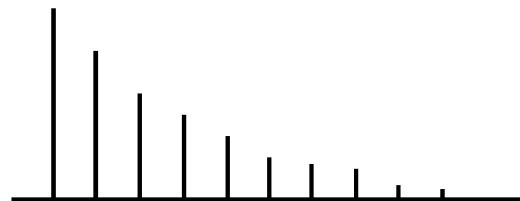
- The *gamma process* can be thought of as the infinitesimal limit of a sequence of gamma random variables.
- Alternatively,
 - if $D \sim \text{DP}(\alpha, H)$
 - and $\gamma \sim \text{Gamma}(\alpha, 1)$
 - then $G = \gamma D \sim \text{GaP}(\alpha H)$
- The gamma distribution is conjugate to the Poisson distribution.





The infinite gamma-Poisson process

- We can associate each atom v_k of the gamma process with a column of a matrix (just like we did with the atoms of a beta process)
- We can generate entries for the matrix as $Z_{nk} \sim \text{Poisson}(v_k)$



IBP

5	4	2	2	1	0	0	1	0
4	4	3	2	0	2	1	0	0
6	2	3	4	0	0	2	0	0
3	5	1	0	3	1	0	1	0
5	3	4	1	1	2	0	0	0
4	4	2	2	2	0	1	0	0

infinite gamma-Poisson





The infinite gamma-Poisson process

- Predictive distribution for the n^{th} row:
 - For each existing feature, sample a count $z_{nk} \sim \text{NegBinom}(m_k, n/(n+1))$

4	2	4	7	0	0	0	0	0
5	0	2	9	4	1	0	0	0
3	2	1	6	2	1	0	0	0
7	1	3	6	3	0	0	0	0





The infinite gamma-Poisson process

- Predictive distribution for the n^{th} row:
 - For each existing feature, sample a count $z_{nk} \sim \text{NegBinom}(m_k, n/(n+1))$

4	2	4	7	0	0	0	0	0
5	0	2	9	4	1	0	0	0
3	2	1	6	2	1	0	0	0
7	1	3	6	3	0	0	0	0
5								





The infinite gamma-Poisson process

- Predictive distribution for the n^{th} row:
 - For each existing feature, sample a count $z_{nk} \sim \text{NegBinom}(m_k, n/(n+1))$

4	2	4	7	0	0	0	0	0
5	0	2	9	4	1	0	0	0
3	2	1	6	2	1	0	0	0
7	1	3	6	3	0	0	0	0
5	0							





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5	0	2	9	4	1	0	0	0
3	2	1	6	2	1	0	0	0
7	1	3	6	3	0	0	0	0
5	0	4	5	2	0			





The infinite gamma-Poisson process

- Predictive distribution for the n^{th} row:
 - For each existing feature, sample a count $z_{nk} \sim \text{NegBinom}(m_k, n/(n+1))$
 - Sample $K_n^* \sim \text{NegBinom}(a, n/(n+1))$

4	2	4	7	0	0	0	0	0
5	0	2	9	4	1	0	0	0
3	2	1	6	2	1	0	0	0
7	1	3	6	3	0	0	0	0
5	0	4	5	2	0			

4





The infinite gamma-Poisson process

- Predictive distribution for the n^{th} row:
 - For each existing feature, sample a count $z_{nk} \sim \text{NegBinom}(m_k, n/(n+1))$.
 - Sample $K_n^* \sim \text{NegBinom}(a, n/(n+1))$.
 - Partition K_n^* according to the CRP, and assign the resulting counts to new columns.

4	2	4	7	0	0	0	0	0
5	0	2	9	4	1	0	0	0
3	2	1	6	2	1	0	0	0
7	1	3	6	3	0	0	0	0
5	0	4	5	2	0	3	1	0

