Distributional Semantic Models

Part 2: The parameters of a DSM

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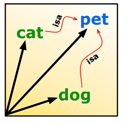
http://wordspace.collocations.de/doku.php/course:start



Outline

General definition of DSMs

A **distributional semantic model** (DSM) is a scaled and/or transformed co-occurrence matrix **M**, such that each row **x** represents the distribution of a target term across contexts.

	get	see	use	hear	eat	kill
knife	0.027	-0.024	0.206	-0.022	-0.044	-0.042
cat	0.031	0.143	-0.243	-0.015	-0.009	0.131
dog	-0.026	0.021	-0.212	0.064	0.013	0.014
boat	-0.022	0.009	-0.044	-0.040	-0.074	-0.042
cup	-0.014	-0.173	-0.249	-0.099	-0.119	-0.042
pig	-0.069	0.094	-0.158	0.000	0.094	0.265
banana	0.047	-0.139	-0.104	-0.022	0.267	-0.042

Term = word, lemma, phrase, morpheme, word pair, ...

General definition of DSMs

Mathematical notation:
- $k \times n$ co-occurrence matrix $\mathbf{M} \in \mathbb{R}^{k \times n}$ (example: 7×6)
  - k rows = **target** terms
  - n columns = **features** or **dimensions**

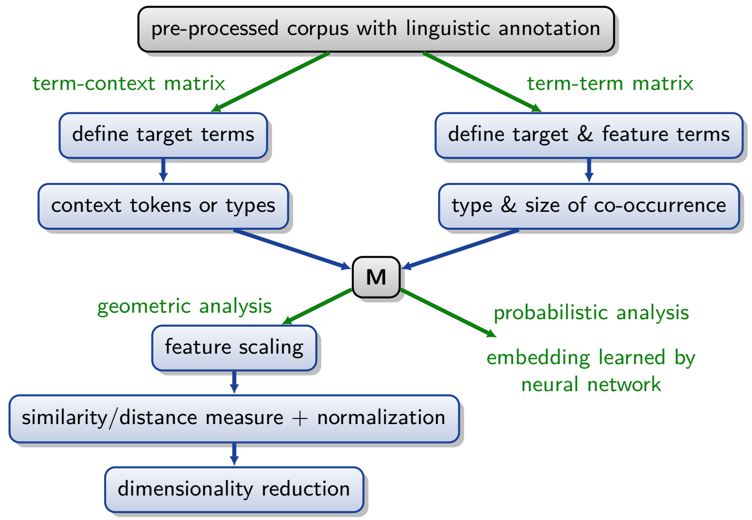
$$\mathbf{M} = \begin{bmatrix} m_{11} & m_{12} & \cdots & m_{1n} \\ m_{21} & m_{22} & \cdots & m_{2n} \\ \vdots & \vdots & & \vdots \\ m_{k1} & m_{k2} & \cdots & m_{kn} \end{bmatrix}$$

- distribution vector $\mathbf{m}_i$ = i-th row of $\mathbf{M}$, e.g. $\mathbf{m}_3 = \mathbf{m}_{\text{dog}} \in \mathbb{R}^n$
- components $\mathbf{m}_i = (m_{i1}, m_{i2}, \ldots, m_{in})$ = features of i-th term:

$$\begin{aligned} \mathbf{m}_3 &= (-0.026, 0.021, -0.212, 0.064, 0.013, 0.014) \\ &= (m_{31}, m_{32}, m_{33}, m_{34}, m_{35}, m_{36}) \end{aligned}$$

Outline

Overview of DSM parameters



Term-context matrix

Term-context matrix records frequency of term in each individual context (e.g. sentence, document, Web page, encyclopaedia article)

$$\mathbf{F} = \begin{bmatrix} \cdots & \mathbf{f}_1 & \cdots \\ \cdots & \mathbf{f}_2 & \cdots \\ & \vdots & \\ & \vdots & \\ \cdots & \mathbf{f}_k & \cdots \end{bmatrix}$$

	Felidae	Pet	Feral	Bloat	Philosophy	Kant	Back pain
cat	10	10	7	–	–	–	–
dog	–	10	4	11	–	–	–
animal	2	15	10	2	–	–	–
time	1	–	–	–	2	1	–
reason	–	1	–	–	1	4	1
cause	–	–	–	2	1	2	6
effect	–	–	–	1	–	1	–

```
> TC <- DSM_TermContext
> head(TC, Inf) # extract full co-oc matrix from DSM object
```


Term-term matrix

Term-term matrix records co-occurrence frequencies with feature terms for each target term

$$\mathbf{M} = \begin{bmatrix} \cdots & \mathbf{m}_1 & \cdots \\ \cdots & \mathbf{m}_2 & \cdots \\ & \vdots & \\ & \vdots & \\ \cdots & \mathbf{m}_k & \cdots \end{bmatrix}$$

	breed	tail	feed	kill	important	explain	likely
cat	83	17	7	37	–	1	–
dog	561	13	30	60	1	2	4
animal	42	10	109	134	13	5	5
time	19	9	29	117	81	34	109
reason	1	–	2	14	68	140	47
cause	–	1	–	4	55	34	55
effect	–	–	1	6	60	35	17

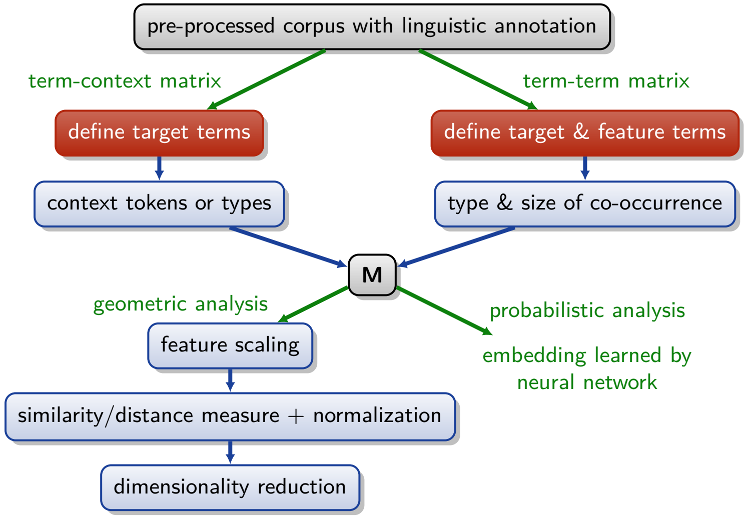
```
> TT <- DSM_TermTerm
> head(TT, Inf)
```

Term-term matrix

Some footnotes:

- Often target terms $\neq$ feature terms
  - e.g. nouns described by co-occurrences with verbs as features
  - identical sets of target & feature terms ➜ symmetric matrix
- Different types of co-occurrence (Evert 2008)
  - **surface context** (word or character window)
  - **textual context** (non-overlapping segments)
  - **syntactic context** (dependency relation)
- Can be seen as smoothing of term-context matrix
  - average over similar contexts (with same context terms)
  - data sparseness reduced, except for small windows
  - we will take a closer look at the relation between term-context and term-term models in part 5 of this tutorial

Overview of DSM parameters



Definition of target and feature terms

- Choice of linguistic unit
  - words
  - bigrams, trigrams, ...
  - multiword units, named entities, phrases, ...
  - morphemes
  - word pairs (☞ analogy tasks)
- Linguistic annotation
  - word forms (minimally requires tokenisation)
  - often lemmatisation or stemming to reduce data sparseness: *go, goes, went, gone, going* ➜ *go*
  - POS disambiguation (*light*/N vs. *light*/A vs. *light*/V)
  - word sense disambiguation (*bank*$_{\text{river}}$ vs. *bank*$_{\text{finance}}$)
  - abstraction: POS tags (or bigrams) as feature terms
- Trade-off between deeper linguistic analysis and
  - need for language-specific resources
  - possible errors introduced at each stage of the analysis

Effects of linguistic annotation

Nearest neighbours of *walk* (BNC)

word forms	lemmatised + POS
stroll	hurry
walking	stroll
walked	stride
go	trudge
path	amble
drive	wander
ride	walk (noun)
wander	walking
sprinted	retrace
sauntered	scuttle

http://clic.cimec.unitn.it/infomap-query/

Effects of linguistic annotation

Nearest neighbours of *arrivare* (Repubblica)

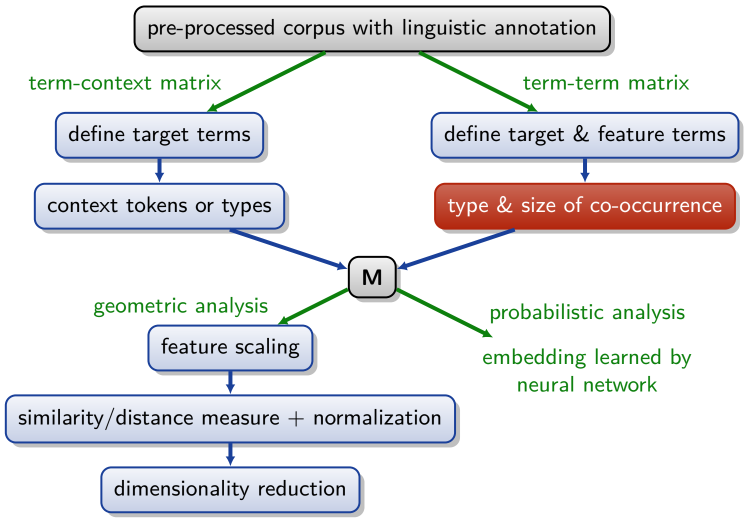
word forms	lemmatised + POS
giungere	giungere
raggiungere	aspettare
arrivi	attendere
raggiungimento	arrivo (noun)
raggiunto	ricevere
trovare	accontentare
raggiunge	approdare
arrivasse	pervenire
arriverà	venire
concludere	piombare

http://clic.cimec.unitn.it/infomap-query/

Selection of target and feature terms

- Full-vocabulary models are often unmanageable
  - 762,424 distinct word forms in BNC, 605,910 lemmata
  - large Web corpora have > 10 million distinct word forms
  - low-frequency targets (and features) do not provide reliable distributional information (too much "noise")
- Frequency-based selection
  - minimum corpus frequency: $f \geq F_{\text{min}}$
  - or accept n_w most frequent terms
  - sometimes also upper threshold: $F_{\text{min}} \leq f \leq F_{\text{max}}$
- Relevance-based selection
  - criterion from IR: document frequency *df*
  - terms with high *df* are too general ➜ uninformative
  - terms with very low *df* may be too sparse to be useful
- Other criteria
  - POS-based filter: no function words, only verbs, . . .

Overview of DSM parameters



Surface context

Context term occurs within a span of *k* words around target.

The silhouette of the sun beyond a wide-open bay on the lake; the sun still glitters although evening has arrived in Kuhmo. It's midsummer; the living room has its instruments and other objects in each of its corners. [L3/R3 span, $k = 6$]

Parameters:

- span size (in words or characters)
- symmetric vs. one-sided span
- uniform or "triangular" (distance-based) weighting
- spans clamped to sentences or other textual units?

Effect of span size

Nearest neighbours of *dog* (BNC)

2-word span	30-word span
cat	kennel
horse	puppy
fox	pet
pet	bitch
rabbit	terrier
pig	rottweiler
animal	canine
mongrel	cat
sheep	to bark
pigeon	Alsatian

http://clic.cimec.unitn.it/infomap-query/

Textual context

Context term is in the same linguistic unit as target.

The silhouette of the sun beyond a wide-open bay on the lake; the sun still glitters although evening has arrived in Kuhmo. It's midsummer; the living room has its instruments and other objects in each of its corners.

Parameters:

- type of linguistic unit
  - sentence
  - paragraph
  - turn in a conversation
  - Web page

Syntactic context

Context term is linked to target by a syntactic dependency (e.g. subject, modifier, . . .).

The silhouette of the sun beyond a wide-open bay on the lake; the sun still glitters although evening has arrived in Kuhmo. It's midsummer; the living room has its instruments and other objects in each of its corners.

Parameters:

- types of syntactic dependency (Padó and Lapata 2007)
- direct vs. indirect dependency paths
  - direct dependencies
  - direct + indirect dependencies
- homogeneous data (e.g. only verb-object) vs. heterogeneous data (e.g. all children and parents of the verb)
- maximal length of dependency path

"Knowledge pattern" context

Context term is linked to target by a lexico-syntactic pattern (text mining, cf. Hearst 1992, Pantel & Pennacchiotti 2008, etc.).

In Provence, Van Gogh painted with bright colors such as red and yellow. These colors produce incredible effects on anybody looking at his paintings.

Parameters:

- inventory of lexical patterns
  - lots of research to identify semantically interesting patterns (cf. Almuhareb & Poesio 2004, Veale & Hao 2008, etc.)
- fixed vs. flexible patterns
  - patterns are mined from large corpora and automatically generalised (optional elements, POS tags or semantic classes)

Structured vs. unstructured context

- In **unstructered** models, context specification acts as a **filter**
  - determines whether context token counts as co-occurrence
  - e.g. muste be linked by any syntactic dependency relation
- In **structured** models, feature terms are **subtyped**
  - depending on their position in the context
  - e.g. left vs. right context, type of syntactic relation, etc.

Structured vs. unstructured surface context

A dog bites a man. The man's dog bites a dog. A dog bites a man.

unstructured	bite
dog	4
man	3

A dog bites a man. The man's dog bites a dog. A dog bites a man.

structured	bite-l	bite-r
dog	3	1
man	1	2

Structured vs. unstructured dependency context

A dog bites a man. The man's dog bites a dog. A dog bites a man.

unstructured	bite
dog	4
man	2

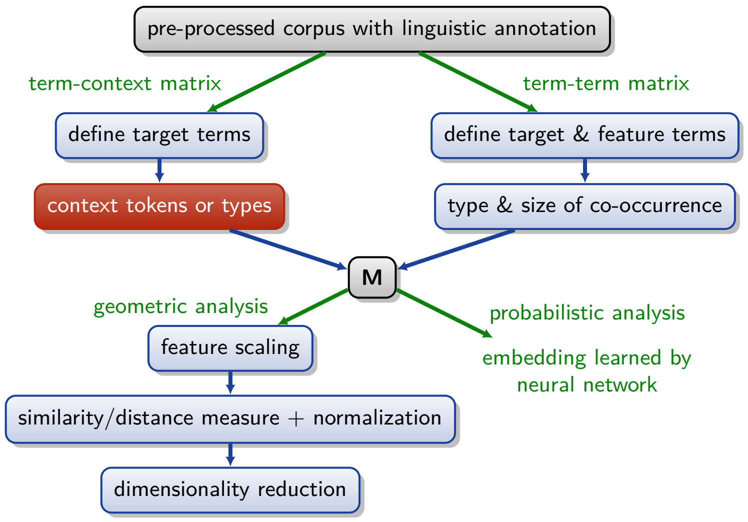
A dog bites a man. The man's dog bites a dog. A dog bites a man.

structured	bite-subj	bite-obj
dog	3	1
man	0	2

Comparison

- Unstructured context
  - data less sparse (e.g. *man kills* and *kills man* both map to the *kill* dimension of the vector $\mathbf{x}_{\text{man}}$)
- Structured context
  - more sensitive to semantic distinctions (*kill-subj* and *kill-obj* are rather different things!)
  - dependency relations provide a form of syntactic "typing" of the DSM dimensions (the "subject" dimensions, the "recipient" dimensions, etc.)
  - important to account for word-order and compositionality

Overview of DSM parameters

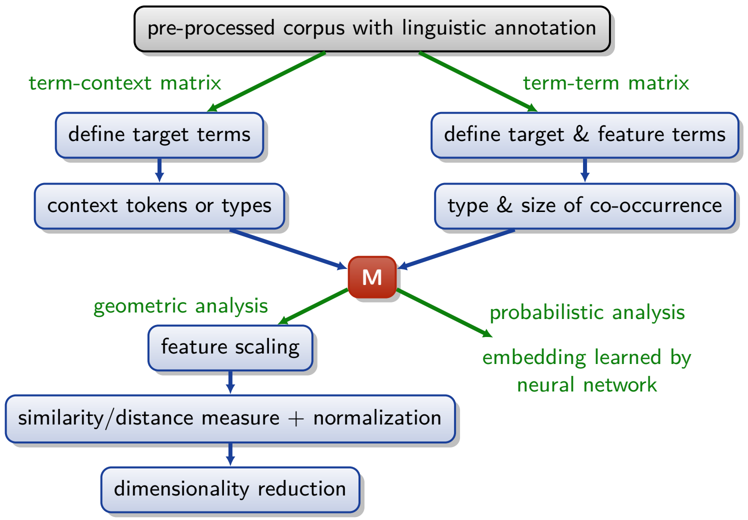


Context tokens vs. context types

- Features are usually context **tokens**, i.e. individual instances
  - document, Wikipedia article, Web page, ...
  - paragraph, sentence, tweet, ...
  - "co-occurrence" count = frequency of term in context token
- Can also be generalised to context **types**, e.g.
  - type = cluster of near-duplicate documents
  - type = syntactic structure of sentence (ignoring content)
  - type = tweets from same author
  - frequency counts from all instances of type are aggregated
- Context types may be anchored at individual tokens
  - n-gram of words (or POS tags) around target
  - subcategorisation pattern of target verb
  - ➥ overlaps with (generalisation of) syntactic co-occurrence

DSM parameters | A taxonomy of DSM parameters

Overview of DSM parameters



Marginal and expected frequencies

- Matrix of observed co-occurrence frequencies not sufficient

target	feature	O	R	C	E
dog	*small*	855	33,338	490,580	134.34
dog	*domesticated*	29	33,338	918	0.25

- Notation
  - O = observed co-occurrence frequency
  - R = overall frequency of target term = row marginal frequency
  - C = overall frequency of feature = column marginal frequency
  - N = sample size $\approx$ size of corpus
- **Expected** co-occurrence **frequency**

$$E = \frac{R \cdot C}{N} \quad \longleftrightarrow \quad O$$

Obtaining marginal frequencies

- Term-document matrix
  - R = frequency of target term in corpus
  - C = size of document (# tokens)
  - N = corpus size
- Syntactic co-occurrence
  - # of dependency instances in which target/feature participates
  - N = total number of dependency instances
  - can be computed from full co-occurrence matrix $\mathbf{M}$
- Textual co-occurrence
  - R, C, O are "document" frequencies, i.e. number of context units in which target, feature or combination occurs
  - N = total # of context units

Obtaining marginal frequencies

- Surface co-occurrence
  - it is quite tricky to obtain fully consistent counts (Evert 2008)
  - at least correct E for span size k (= number of tokens in span)

$$E = k \cdot \frac{R \cdot C}{N}$$

  with R, C = individual corpus frequencies and N = corpus size
  - can also be implemented by pre-multiplying $R' = k \cdot R$
  - ☞ alternatively, compute marginals and sample size by summing over full co-occurrence matrix (➜ E as above, but inflated N)
- NB: shifted PPMI (Levy and Goldberg 2014) corresponds to a post-hoc application of the span size adjustment
  - performs worse than PPMI, but paper suggests they already approximate correct E by summing over co-occurrence matrix

Marginal frequencies in `wordspace`

DSM objects in `wordspace` include marginal frequencies as well as counts of nonzero cells for rows and columns.

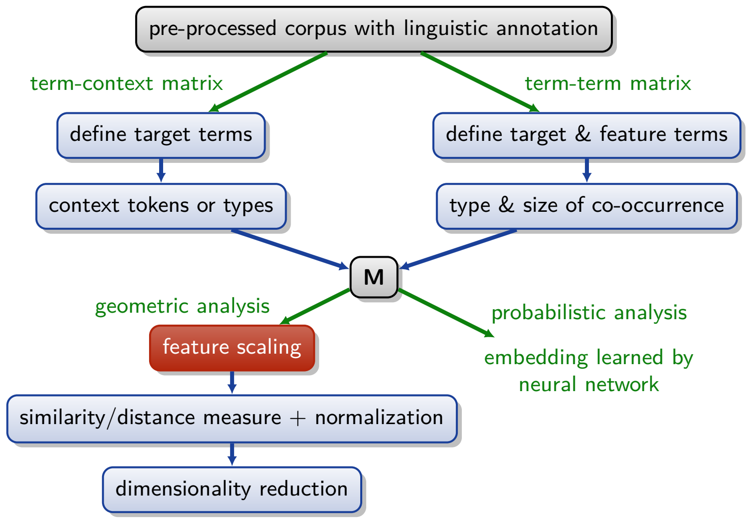
```
> TT$rows
    term       f nnzero
1    cat   22007      5
2    dog   50807      7
3 animal   77053      7
4   time 1156693      7
5 reason   95047      6
6  cause   54739      5
7 effect  133102      6
> TT$cols
...
> TT$globals$N
[1] 199902178
> TT$M # the full co-occurrence matrix
```


Geometric vs. probabilistic interpretation

- Geometric interpretation
  - row vectors as points or arrows in n-dimensional space
  - very intuitive, good for visualisation
  - use techniques from geometry and matrix algebra
- Probabilistic interpretation
  - co-occurrence matrix as observed sample statistic that is "explained" by a generative probabilistic model
  - e.g. probabilistic LSA (Hoffmann 1999), Latent Semantic Clustering (Rooth *et al.* 1999), Latent Dirichlet Allocation (Blei *et al.* 2003), etc.
  - explicitly accounts for random variation of frequency counts
  - recent work: neural word embeddings

☞ focus on geometric interpretation in this tutorial

Overview of DSM parameters



Feature scaling

Feature scaling is used to "discount" less important features:

- Logarithmic scaling: $O' = \log(O + 1)$
  (cf. Weber-Fechner law for human perception)
- Relevance weighting, e.g. tf.idf (information retrieval)

$$tf.idf = tf \cdot \log(D/df)$$

  - tf = co-occurrence frequency O
  - df = document frequency of feature (or nonzero count)
  - D = total number of documents (or row count of $\mathbf{M}$)
- Statistical **association measures** (Evert 2004, 2008) take frequency of target term and feature into account
  - often based on comparison of observed and expected co-occurrence frequency
  - measures differ in how they balance O and E

Simple association measures

- pointwise Mutual Information (MI)

$$\mathrm{MI} = \log_2 \frac{O}{E}$$

- local MI

$$\text{local-MI} = O \cdot \mathrm{MI} = O \cdot \log_2 \frac{O}{E}$$

- t-score

$$t = \frac{O - E}{\sqrt{O}}$$

target	feature	O	E	MI	local-MI	t-score
dog	*small*	855	134.34	2.67	2282.88	24.64
dog	*domesticated*	29	0.25	6.85	198.76	5.34
dog	*sgjkj*	1	0.00027	11.85	11.85	1.00

Other association measures

- simple log-likelihood (≈ local-MI)

$$G^2 = \pm 2 \cdot \left(O \cdot \log_2 \frac{O}{E} - (O - E) \right)$$

  with positive sign for $O > E$ and negative sign for $O < E$
- Dice coefficient

$$\mathrm{Dice} = \frac{2O}{R + C}$$

- Many other simple association measures (AMs) available
- Further AMs computed from full contingency tables, see
  - Evert (2008)
  - `http://www.collocations.de/`
  - `http://sigil.r-forge.r-project.org/`

Applying association scores in wordspace

```
> options(digits=3)  # print fractional values with limited precision
> dsm.score(TT, score="MI", sparse=FALSE, matrix=TRUE)
        breed   tail   feed   kill important explain  likely
cat      6.21  4.568  3.129  2.801      -Inf  0.0182    -Inf
dog      7.78  3.081  3.922  2.323    -3.774 -1.1888 -0.4958
animal   3.50  2.132  4.747  2.832    -0.674 -0.4677 -0.0966
time    -1.65 -2.236 -0.729 -1.097    -1.728 -1.2382  0.6392
reason  -2.30   -Inf -1.982 -0.388     1.472  4.0368  2.8860
cause    -Inf -0.834   -Inf -2.177     1.900  2.8329  4.0691
effect   -Inf -2.116 -2.468 -2.459     0.791  1.6312  0.9221
```

☞ sparseness of the matrix has been lost!

☞ cells with score $x = -\infty$ are inconvenient

☞ distribution of scores may be even more skewed than co-occurrence frequencies (esp. for local-MI)

Sparse association measures

- Sparse association scores are cut off at zero, i.e.

$$f(x) = \begin{cases} x & x > 0 \\ 0 & x \leq 0 \end{cases}$$

- Also known as "positive" scores
  - PPMI = positive pointwise MI (e.g. Bullinaria and Levy 2007)
  - `wordspace` computes sparse AMs by default ➜ `"MI"` = PPMI
- Preserves sparseness if $x \leq 0$ for all empty cells ($O = 0$)
  - sparseness may even increase: cells with $x < 0$ become empty
- Usually combined with signed association measure satisfying
  - $x > 0$ for $O > E$
  - $x < 0$ for $O < E$

Score transformations

An additional scale transformation can be applied in order to de-skew association scores:

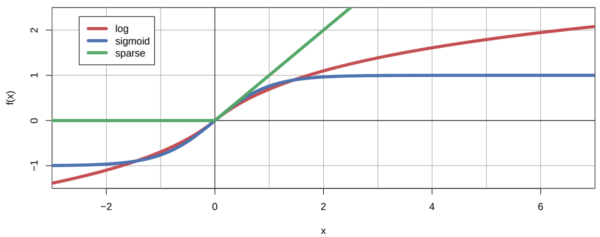
- signed logarithmic transformation

$$f(x) = \pm \log(|x| + 1)$$

- sigmoid transformation as soft binarization

$$f(x) = \tanh x$$

- sparse AM as cutoff transformation



Association scores & transformations in wordspace

```
> dsm.score(TT, score="MI", matrix=TRUE)  # PPMI
       breed tail feed kill important explain likely
cat     6.21 4.57 3.13 2.80     0.000  0.0182  0.000
dog     7.78 3.08 3.92 2.32     0.000  0.0000  0.000
animal  3.50 2.13 4.75 2.83     0.000  0.0000  0.000
time    0.00 0.00 0.00 0.00     0.000  0.0000  0.639
reason  0.00 0.00 0.00 0.00     1.472  4.0368  2.886
cause   0.00 0.00 0.00 0.00     1.900  2.8329  4.069
effect  0.00 0.00 0.00 0.00     0.791  1.6312  0.922
> dsm.score(TT, score="simple-ll", matrix=TRUE)
> dsm.score(TT, score="simple-ll", transf="log", matrix=T)
# logarithmic co-occurrence frequency
> dsm.score(TT, score="freq", transform="log", matrix=T)

# now try other parameter combinations
> ?dsm.score  # read help page for available parameter settings
```

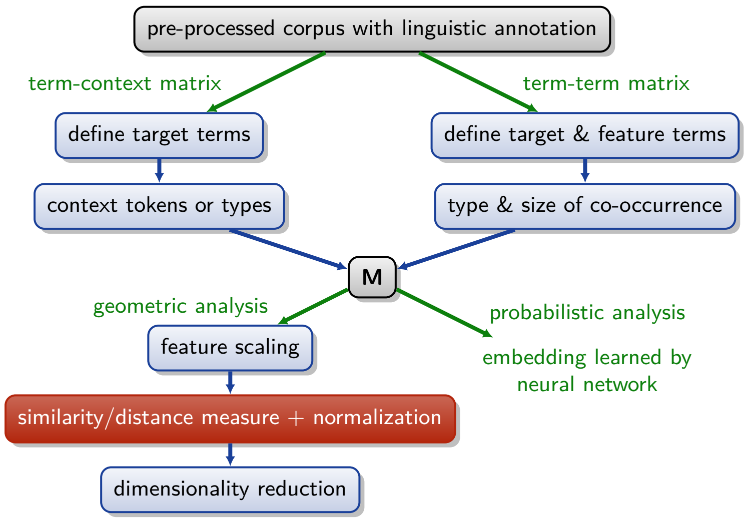
Scaling of column vectors

- In statistical analysis and machine learning, features are usually centred and scaled so that

$$\text{mean} \quad \mu = 0$$
$$\text{variance} \quad \sigma^2 = 1$$

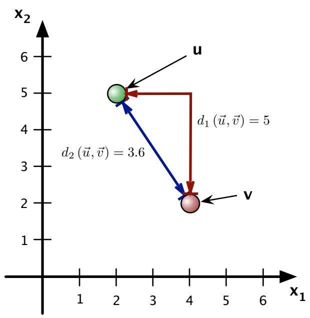
- In DSM research, this step is less common for columns of **M**
  - centring is a prerequisite for certain dimensionality reduction and data analysis techniques (esp. PCA)
  - but co-occurrence matrix no longer sparse!
  - scaling may give too much weight to rare features
- **M** cannot be row-normalised and column-scaled at the same time (result depends on ordering of the two steps)

Overview of DSM parameters



Geometric distance = metric

- **Distance** between vectors $\mathbf{u}, \mathbf{v} \in \mathbb{R}^n$ ➜ (dis)similarity
  - $\mathbf{u} = (u_1, \dots, u_n)$
  - $\mathbf{v} = (v_1, \dots, v_n)$
- **Euclidean** distance $d_2(\mathbf{u}, \mathbf{v})$
- "City block" **Manhattan** distance $d_1(\mathbf{u}, \mathbf{v})$
- Both are special cases of the **Minkowski** p-distance $d_p(\mathbf{u}, \mathbf{v})$ (for $p \in [1, \infty]$)

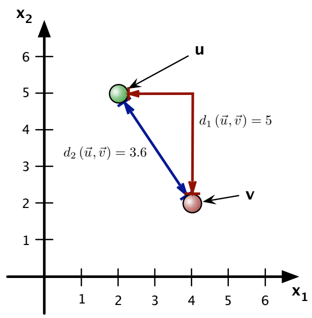


$$d_p(\mathbf{u}, \mathbf{v}) := \left(|u_1 - v_1|^p + \dots + |u_n - v_n|^p\right)^{1/p}$$

$$d_\infty(\mathbf{u}, \mathbf{v}) = \max\{|u_1 - v_1|, \dots, |u_n - v_n|\}$$

Geometric distance = metric

- **Distance** between vectors $\mathbf{u}, \mathbf{v} \in \mathbb{R}^n$ ➜ (dis)similarity
  - $\mathbf{u} = (u_1, \dots, u_n)$
  - $\mathbf{v} = (v_1, \dots, v_n)$
- **Euclidean** distance $d_2(\mathbf{u}, \mathbf{v})$
- "City block" **Manhattan** distance $d_1(\mathbf{u}, \mathbf{v})$
- Extension of p-distance $d_p(\mathbf{u}, \mathbf{v})$ (for $0 \le p \le 1$)



$$d_p(\mathbf{u}, \mathbf{v}) := |u_1 - v_1|^p + \dots + |u_n - v_n|^p$$

$$d_0(\mathbf{u}, \mathbf{v}) = \#\{i \mid u_i \neq v_i\}$$

Computing distances

Preparation: store "scored" matrix in DSM object

```
> TT <- dsm.score(TT, score="freq", transform="log")
```

Compute distances between individual term pairs …

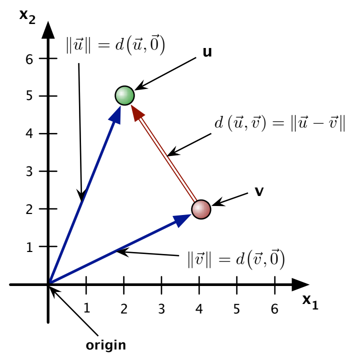
```
> pair.distances(c("cat","cause"), c("animal","effect"),
                 TT, method="euclidean")
 cat/animal cause/effect
       4.16         1.53
```

… or full distance matrix.

```
> dist.matrix(TT, method="euclidean")
> dist.matrix(TT, method="minkowski", p=4)
```


Distance and vector length = norm

- Intuitively, distance $d(\mathbf{u},\mathbf{v})$ should correspond to length $\|\mathbf{u}-\mathbf{v}\|$ of displacement vector $\mathbf{u}-\mathbf{v}$
  - $d(\mathbf{u},\mathbf{v})$ is a **metric**
  - $\|\mathbf{u}-\mathbf{v}\|$ is a **norm**
  - $\|\mathbf{u}\| = d(\mathbf{u},\mathbf{0})$
- Such a metric is always **translation-invariant**

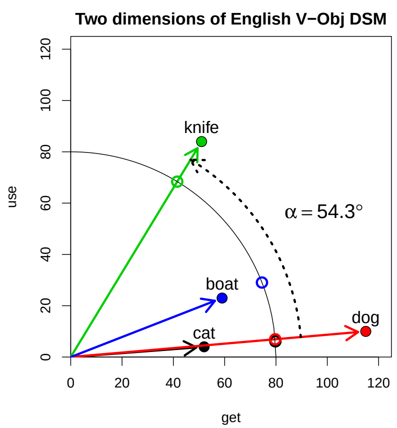


- $d_p(\mathbf{u},\mathbf{v}) = \|\mathbf{u}-\mathbf{v}\|_p$
- **Minkowski p-norm** for $p \in [1,\infty]$ (not $p < 1$):

$$\|\mathbf{u}\|_p := \left(|u_1|^p + \cdots + |u_n|^p\right)^{1/p}$$

Normalisation of row vectors

- Geometric distances only meaningful for vectors of the same length $\|\mathbf{x}\|$
- Normalize by scalar division: $\mathbf{x}' = \mathbf{x}/\|\mathbf{x}\| = \left(\frac{x_1}{\|\mathbf{x}\|}, \frac{x_2}{\|\mathbf{x}\|}, \ldots\right)$ with $\|\mathbf{x}'\| = 1$
- Norm must be compatible with distance measure!
- Special case: scale to relative frequencies with $\|\mathbf{x}\|_1 = |x_1| + \cdots + |x_n|$
  ➜ probabilistic interpretation



Norms and normalization

```
> rowNorms(TT$S, method="euclidean")
  cat   dog animal   time reason  cause effect
 6.90  8.96   8.82  10.29   8.13   6.86   6.52
```

```
> TT <- dsm.score(TT, score="freq", transform="log",
                  normalize=TRUE, method="euclidean")
> rowNorms(TT$S, method="euclidean") # all = 1 now
> dist.matrix(TT, method="euclidean")
         cat   dog animal  time reason cause effect
cat    0.000 0.224  0.473 0.782  1.121 1.239  1.161
dog    0.224 0.000  0.398 0.698  1.065 1.179  1.113
animal 0.473 0.398  0.000 0.426  0.841 0.971  0.860
time   0.782 0.698  0.426 0.000  0.475 0.585  0.502
reason 1.121 1.065  0.841 0.475  0.000 0.277  0.198
cause  1.239 1.179  0.971 0.585  0.277 0.000  0.224
effect 1.161 1.113  0.860 0.502  0.198 0.224  0.000
```

Other distance measures

- Information theory: **Kullback-Leibler** (KL) **divergence** for probability vectors (☞ non-negative, $\|\mathbf{x}\|_1 = 1$)

$$D(\mathbf{u}\|\mathbf{v}) = \sum_{i=1}^{n} u_i \cdot \log_2 \frac{u_i}{v_i}$$

- Properties of KL divergence
  - most appropriate in a probabilistic interpretation of $\mathbf{M}$
  - zeroes in $\mathbf{v}$ without corresponding zeroes in $\mathbf{u}$ are problematic
  - not symmetric, unlike geometric distance measures
  - alternatives: skew divergence, Jensen-Shannon divergence
- A symmetric distance measure (Endres and Schindelin 2003)

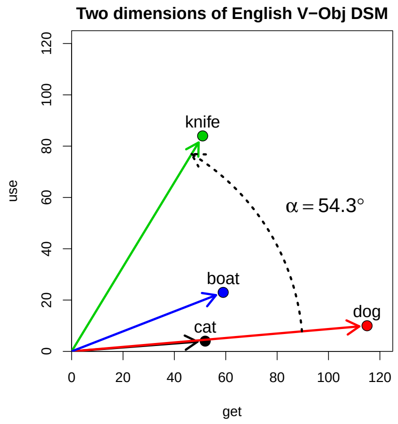
$$D_{\mathbf{uv}} = D(\mathbf{u}\|\mathbf{z}) + D(\mathbf{v}\|\mathbf{z}) \quad \text{with} \quad \mathbf{z} = \frac{\mathbf{u}+\mathbf{v}}{2}$$

Similarity measures

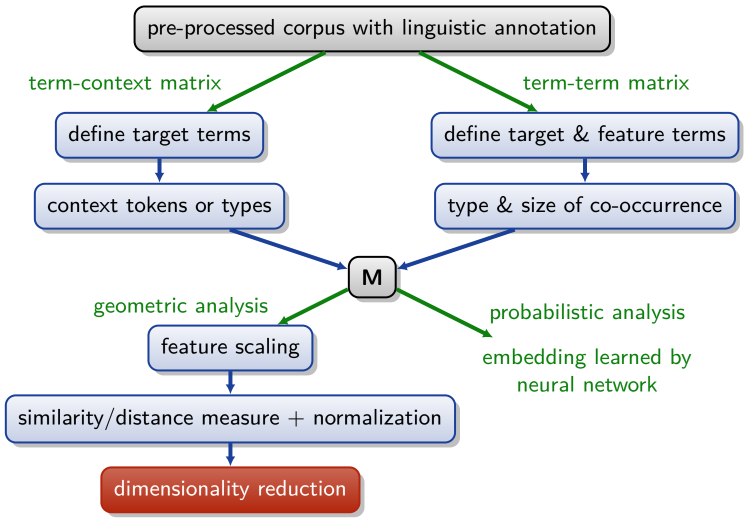
- Angle α between vectors $\mathbf{u}, \mathbf{v} \in \mathbb{R}^n$ is given by

$$\cos\alpha = \frac{\sum_{i=1}^{n} u_i \cdot v_i}{\sqrt{\sum_i u_i^2} \cdot \sqrt{\sum_i v_i^2}} = \frac{\mathbf{u}^T\mathbf{v}}{\|\mathbf{u}\|_2 \cdot \|\mathbf{v}\|_2}$$

- **cosine** measure of similarity: $\cos\alpha$
  - $\cos\alpha = 1$ ➜ collinear
  - $\cos\alpha = 0$ ➜ orthogonal
- Corresponding metric: angular distance α



Overview of DSM parameters



Dimensionality reduction = model compression

- Co-occurrence matrix $\mathbf{M}$ is often unmanageably large and can be extremely sparse
  - Google Web1T5: 1M × 1M matrix with one trillion cells, of which less than 0.05% contain nonzero counts (Evert 2010)

➥ Compress matrix by reducing dimensionality (= rows)

- **Feature selection**: columns with high frequency & variance
  - measured by entropy, chi-squared test, nonzero count, . . .
  - may select similar dimensions and discard valuable information
  - joint selection of multiple features is useful but expensive
- **Projection** into (linear) subspace
  - principal component analysis (PCA)
  - independent component analysis (ICA)
  - random indexing (RI)
  - ☞ intuition: preserve distances between data points

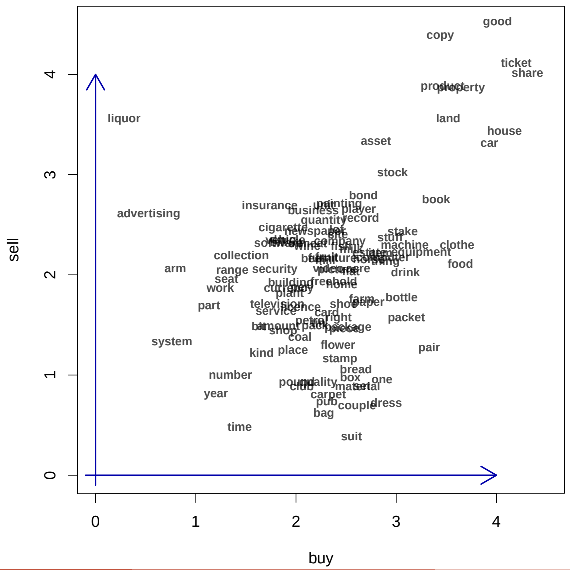
Dimensionality reduction & latent dimensions

Landauer and Dumais (1997) claim that LSA dimensionality reduction (and related PCA technique) uncovers **latent dimensions** by exploiting correlations between features.

- Example: term-term matrix
- V-Obj cooc's extracted from BNC
  - targets = noun lemmas
  - features = verb lemmas
- feature scaling: association scores (modified log Dice coefficient)
- $k = 111$ nouns with $f \geq 20$ (must have non-zero row vectors)
- $n = 2$ dimensions: *buy* and *sell*

noun	*buy*	*sell*
bond	0.28	0.77
cigarette	-0.52	0.44
dress	0.51	-1.30
freehold	-0.01	-0.08
land	1.13	1.54
number	-1.05	-1.02
per	-0.35	-0.16
pub	-0.08	-1.30
share	1.92	1.99
system	-1.63	-0.70

Dimensionality reduction & latent dimensions

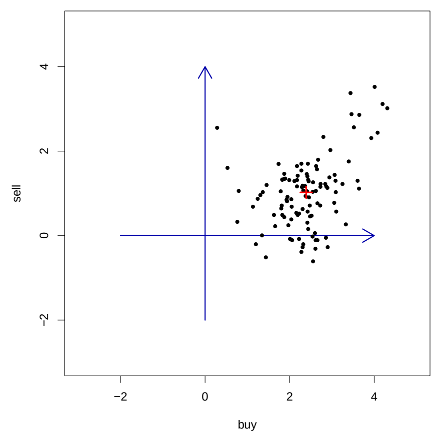


Motivating latent dimensions & subspace projection

- The **latent property** of being a commodity is "expressed" through associations with several verbs: *sell*, *buy*, *acquire*, ...
- Consequence: these DSM dimensions will be **correlated**
- Identify **latent dimension** by looking for strong correlations (or weaker correlations between large sets of features)
- Projection into subspace V of $k < n$ latent dimensions as a "**noise reduction**" technique ➜ **LSA**
- Assumptions of this approach:
  - "latent" distances in V are semantically meaningful
  - other "residual" dimensions represent chance co-occurrence patterns, often particular to the corpus underlying the DSM

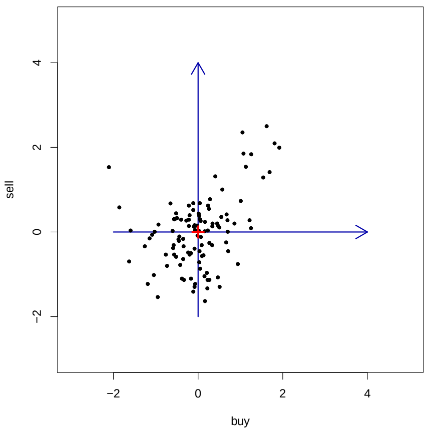
Centering the data set

- **Uncentered data set**
- Centered data set
- Variance of centered data



Centering the data set

- Uncentered data set
- **Centered data set**
- Variance of centered data

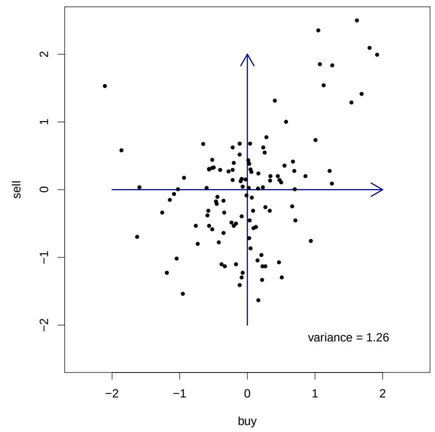


DSM parameters | A taxonomy of DSM parameters

Centering the data set

- Uncentered data set
- Centered data set
- **Variance of centered data**

$$\sigma^2 = \frac{1}{k-1} \sum_{i=1}^{k} \|\mathbf{x}^{(i)}\|^2$$



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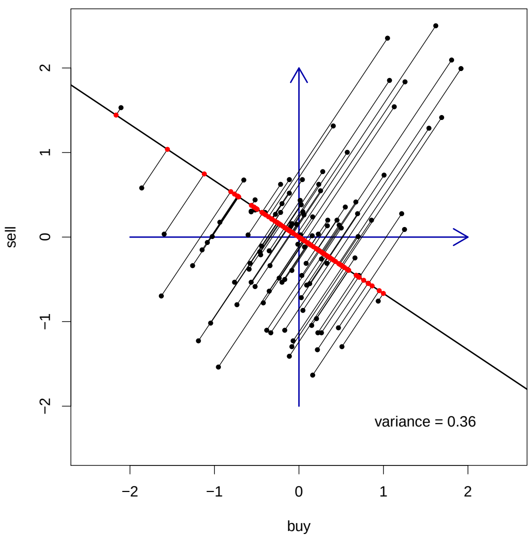
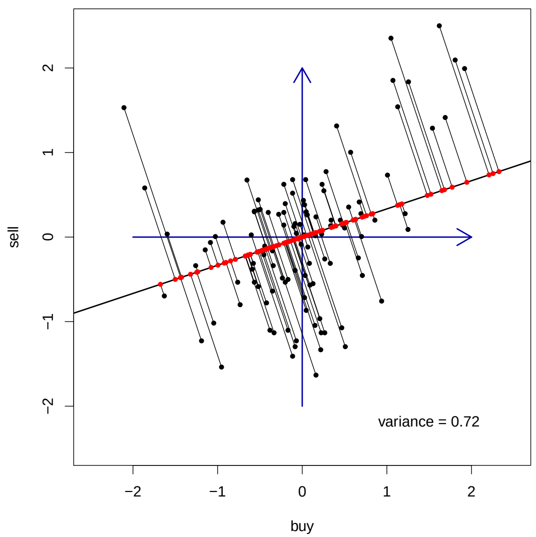
DSM parameters | A taxonomy of DSM parameters

Projection and preserved variance: examples



DSM parameters | A taxonomy of DSM parameters

Projection and preserved variance: examples



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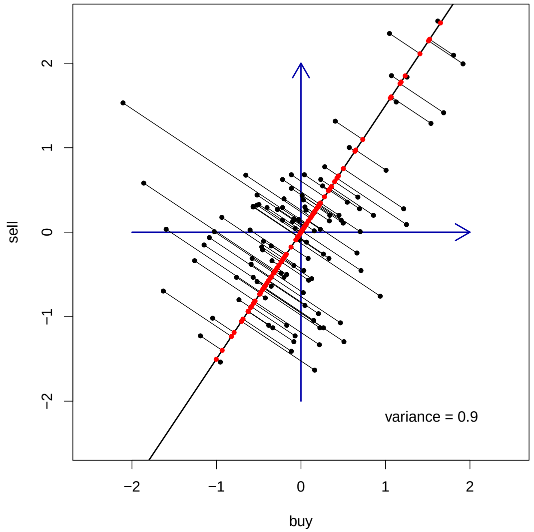
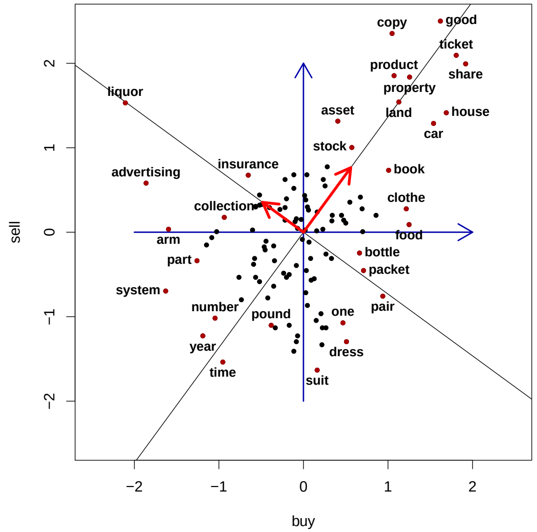
Projection and preserved variance: examples



Orthogonal PCA dimensions



Dimensionality reduction in practice

```
# it is customary to omit the centring: SVD dimensionality reduction
> TT2 <- dsm.projection(TT, n=2, method="svd")
> TT2
         svd1    svd2
cat    -0.733 -0.6615
dog    -0.782 -0.6110
animal -0.914 -0.3606
time   -0.993  0.0302
reason -0.889  0.4339
cause  -0.817  0.5615
effect -0.871  0.4794

> x <- TT2[, 1]  # first latent dimension
> y <- TT2[, 2]  # second latent dimension
> plot(TT2, pch=20, col="red",
       xlim=extendrange(x), ylim=extendrange(y))
> text(TT2, rownames(TT2), pos=3)
```


Outline

Some well-known DSM examples

Latent Semantic Analysis (Landauer and Dumais 1997)

- term-context matrix with document context
- weighting: log term frequency and term entropy
- distance measure: cosine
- dimensionality reduction: SVD

Hyperspace Analogue to Language (Lund and Burgess 1996)

- term-term matrix with surface context
- structured (left/right) and distance-weighted frequency counts
- distance measure: Minkowski metric ($1 \leq p \leq 2$)
- dimensionality reduction: feature selection (high variance)

Some well-known DSM examples

Infomap NLP (Widdows 2004)

- term-term matrix with unstructured surface context
- weighting: none
- distance measure: cosine
- dimensionality reduction: SVD

Random Indexing (Karlgren and Sahlgren 2001)

- term-term matrix with unstructured surface context
- weighting: various methods
- distance measure: various methods
- dimensionality reduction: random indexing (RI)

Some well-known DSM examples

Dependency Vectors (Padó and Lapata 2007)

- term-term matrix with unstructured dependency context
- weighting: log-likelihood ratio
- distance measure: PPMI-weighted Dice (Lin 1998)
- dimensionality reduction: none

Distributional Memory (Baroni and Lenci 2010)

- term-term matrix with structured and unstructered dependencies + knowledge patterns
- weighting: local-MI on type frequencies of link patterns
- distance measure: cosine
- dimensionality reduction: none

Outline

Scaling up to the real world

- So far, we have worked on minuscule **toy models**
- ☞ We want to scale up to **real world** data sets now

- Example 1: window-based DSM on BNC content words
  - 83,926 lemma types with $f \geq 10$
  - term-term matrix with $83{,}926 \cdot 83{,}926 = 7$ billion entries
  - standard representation requires 56 GB of RAM (8-byte floats)
  - only 22.1 million non-zero entries (= 0.32%)

- Example 2: Google Web 1T 5-grams (1 trillion words)
  - more than 1 million word types with $f \geq 2500$
  - term-term matrix with 1 trillion entries requires 8 TB RAM
  - only 400 million non-zero entries (= 0.04%)

Sparse matrix representation

- Invented example of a **sparsely populated** DSM matrix

	eat	get	hear	kill	see	use
boat	·	59	·	·	39	23
cat	·	·	·	26	58	·
cup	·	98	·	·	·	·
dog	33	·	42	·	83	·
knife	·	·	·	·	·	84
pig	9	·	·	27	·	·

- Store only non-zero entries in compact **sparse matrix format**

row	col	value	row	col	value
1	2	59	4	1	33
1	5	39	4	3	42
1	6	23	4	5	83
2	4	26	5	6	84
2	5	58	6	1	9
3	2	98	6	4	27

Working with sparse matrices

- Compressed format: each row index (or column index) stored only once, followed by non-zero entries in this row (or column)
  - convention: **column-major** matrix (data stored by columns)

- Specialised algorithms for sparse matrix algebra
  - especially matrix multiplication, solving linear systems, etc.
  - take care to avoid operations that create a dense matrix!

- **R** implementation: `Matrix` package
  - essential for real-life distributional semantics
  - `wordspace` provides additional support for sparse matrices (vector distances, sparse SVD, ...)

- Other software: Matlab, Octave, Python + SciPy

Outline

Triplet tables

- A sparse DSM matrix can be represented as a table of triplets (target, feature, co-occurrence frequency)
  - for syntactic co-occurrence and term-document matrices, marginals can be computed from a complete triplet table
  - for surface and textual co-occurrence, marginals have to be provided in separate files (see `?read.dsm.triplet`)

noun	rel	verb	f	mode
dog	subj	bite	3	spoken
dog	subj	bite	12	written
dog	obj	bite	4	written
dog	obj	stroke	3	written
...	...	...	...	...

- `DSM_VerbNounTriples_BNC` contains additional information
  - syntactic relation between noun and verb
  - written or spoken part of the British National Corpus

Constructing a DSM from a triplet table

- Additional information can be used for filtering (verb-object relation), or aggregate frequencies (spoken + written BNC)

```
> tri <- subset(DSM_VerbNounTriples_BNC, rel == "obj")
```

- Construct DSM object from triplet input
  - `raw.freq=TRUE` indicates raw co-occurrence frequencies (rather than a pre-weighted DSM)
  - constructor aggregates counts from duplicate entries
  - marginal frequencies are automatically computed

```
> VObj <- dsm(target=tri$noun, feature=tri$verb,
              score=tri$f, raw.freq=TRUE)
> VObj # inspect marginal frequencies (e.g. head(VObj$rows, 20))
```


Exploring the DSM

```
> VObj <- dsm.score(VObj, score="MI", normalize=TRUE)

> nearest.neighbours(VObj, "dog") # angular distance
   horse     cat   animal  rabbit    fish     guy
    73.9    75.9     76.2    77.0    77.2    78.5
 cichlid     kid      bee creature
    78.6    79.0     79.1    79.5

> nearest.neighbours(VObj, "dog", method="manhattan")
# NB: we used an incompatible Euclidean normalization!

> VObj50 <- dsm.projection(VObj, n=50, method="svd")
> nearest.neighbours(VObj50, "dog")
```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