Inter Annotator Agreement Workshop SANTA

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Introduction

Gamma

Combining Expected and Observed Agreement Calculating Expected Agreement Calculating Observed Agreement

Results

All Annotations Own vs. Foreign Annotations Own vs. Student Annotations Foreign vs. Student Annotations Comments

Section 1

Introduction

Motivation

- IAA expresses agreement between annotators/raters quantitatively
- Often used as an upper bound in NLP:
 Computers can't be expected to perform better than human agreement
- Annotations with high IAA are considered more reliable
- Sometimes used to steer guideline/resource development
 - ► '90% solution': Remove word senses for which annotators achieve less than 90%

 Hovy et al. (2006)
- Corpus releases should be accompanied by IAA values, to allow estimation of annotation quality

Different Metrics

- Not all annotation tasks are the same
 - PoS tagging: Assign each word to a category
 - Only categorizing
 - Sentence splitting: Mark sentence boundaries
 - Only unitizing
 - Named entities: Select a span and assign it to a category
 - Unitizing, categorizing
- Different metrics for different tasks!

Cohen 1960; Fleiss 1971; Fournier and Inkpen 2012; Mathet et al. 2015

Different Metrics

Common Properties

- All metrics incorporate observed and expected agreement
- Observed agreement: Extracted from the annotations
- Expected agreement: Agreement to be expected by chance annotations
 - Indicates difficulty of the annotation task
 - Allows comparing agreement values with different numbers of categories!

Expected Agreement

If two annotators assign word classes (noun, verb, adjective, other) by throwing a 4-sided die, they achieve a certain level of agreement (this is a categorization task).



Gamma

Section 2

Gamma

Metric γ has been published in this paper: Yann Mathet et al. "The Unified and Holistic Method Gamma (γ) for Inter-Annotator Agreement Measure and Alignment". In: Computational Linguistics 41.3 (2015), pp. 437–479

Three Components

- Combination of expected and observed agreement
- Calculation of expected agreement
- Calculation of observed agreement

Combining Expected and Observed Agreement

Note: γ is defined based on disagreements! Assuming we have calculated observed (δ_o) and expected (δ_e) disagreement

$$\gamma = 1 - \frac{\delta_o}{\delta_o} \tag{1}$$

Combining Expected and Observed Agreement Examples

$$\gamma = 1 - \frac{\delta_o}{\delta_e}$$

Combining Expected and Observed Agreement

Examples

$$\gamma = 1 - rac{\delta_o}{\delta_e}$$

δ_o	δ_e	γ	
0.99 0.01	0.01 0.99	0.98 -98	(upper bound: 1) (lower bound: $-\infty$)
0.5 0.5 0.5	0.25 0.5 0.75	$-1 \\ 0 \\ 0.33$	(a secondary
0.25 0.5 0.75	0.5 0.5 0.5	$0.5 \\ 0 \\ -0.5$	

Table: γ scores for observed (δ_o) and expected (δ_e) disagreement

Calculating Expected Agreement

- Random annotations need to be realistic w.r.t. several criteria
 - Distribution of units per annotator
 - Distribution of categories
 - **...**
- γ's expected disagreement is based on real annotations
 - 1. Take the annotations created by a real annotator
 - 2. Split the text at a random point
 - 3. Permute the two parts
 - 4. Repeat multiple times and calculate disagreement
- This doesn't work if the text only contains a single annotation that spans the entire text

Basics

- Local level: Measuring dissimilarity between two annotations
- Global level: Create unitary alignments over all annotations by all annotators

Situations

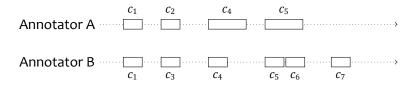


Figure: Two annotators and (some) possible situations

One Annotation is defined by

- begin/end
- feature values (including category)

If these are the same, we consider two annotations to be equal

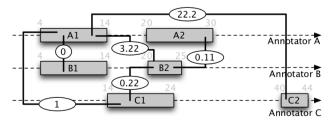
Positional Dissimilarity

$$d_{pos}(u,v) = \left(\underbrace{\frac{|\operatorname{start difference}}{|\operatorname{start}(u) - \operatorname{start}(v)|}_{|\operatorname{length of u}} + \underbrace{|\operatorname{end}(u) - \operatorname{end}(v)|}_{|\operatorname{length of v}} \right)^2$$

Positional Dissimilarity

$$d_{pos}(u,v) = \underbrace{\left(\underbrace{\frac{|\operatorname{start}\operatorname{difference}}{|\operatorname{start}(u) - \operatorname{start}(v)|} + \underbrace{|\operatorname{end}(u) - \operatorname{end}(v)|}_{|\operatorname{end}(v) - \operatorname{start}(v)|} + \underbrace{(\operatorname{end}(v) - \operatorname{start}(v))}_{|\operatorname{ength}\operatorname{of}\operatorname{u}} \right)^{T}}_{|\operatorname{length}\operatorname{of}\operatorname{u}}$$

Examples



Positional Dissimilarity

$$d_{pos}(u,v) = \left(\underbrace{\frac{\left| \operatorname{start} \operatorname{difference} \right|}{\left| \operatorname{start}(u) - \operatorname{start}(v) \right|} + \left| \operatorname{end}(u) - \operatorname{end}(v) \right|}_{\left| \operatorname{end}(u) - \operatorname{start}(v) \right|} + \underbrace{\left(\operatorname{end}(v) - \operatorname{start}(v) \right)}_{\left| \operatorname{ength} \operatorname{of} u \right|}^{2} \right)^{2}}_{\left| \operatorname{ength} \operatorname{of} u \right|}$$

Here: Token numbers as positions (using a heuristic tokenizer).

Categorial Dissimilarity

Gamma

Define dissimilarity between categories in a matrix

	<i>c</i> ₁	c_2	<i>c</i> ₃
c_1	0	0.5	1
c_2	0.5	0	0.25
<i>c</i> ₃	1	0.25	0

Categorial Dissimilarity

Gamma

Define dissimilarity between categories in a matrix

	c_1	c_2	<i>c</i> ₃
c_1	0	0.5	1
c_2	0.5	0	0.25
<i>c</i> ₃	1	0.25	0

SANTA

$$d_{cat}(u, v) = \left\{ egin{array}{ll} 0 & ext{if } \operatorname{cat}(u) = \operatorname{cat}(v) \ 1 & ext{otherwise} \end{array}
ight.$$

I.e.: We don't use graded dissimilarity here

Categorial Dissimilarity: Features vs. Categories

Most guidelines define several individual features, instead of a single category. Feature assignments have been merged into a single string to represent a 'category'.

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Example

Addressee : Mouse Speaker : Mouse

becomes the 'category' Addressee=Mouse+Speaker=Mouse

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Example

Addressee : Mouse Speaker : Mouse

becomes the 'category' Addressee=Mouse+Speaker=Mouse

- This is a shortcoming
- Guideline authors: Define severity of disagreement between categories

Combining Dissimilarity

$$d_{combi}(u, v) = \alpha d_{pos}(u, v) + \beta d_{cat}(u, v)$$

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Intuitions and Remarks

- $ightharpoonup \alpha$ and β can be used to express importance
 - ightharpoonup Our setting, $\alpha = \beta = 1$
 - ▶ I.e., positional and categorial disagreement are equally important

Combining Dissimilarity

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Intuitions and Remarks

- $ightharpoonup \alpha$ and β can be used to express importance
 - ightharpoonup Our setting, $\alpha = \beta = 1$
 - ▶ I.e., positional and categorial disagreement are equally important
- Dissimilarity between two annotations is roughly between 0 (zero) and the squared length of the text (because of the positional dissimilarity)

Alignment

- ▶ Pairwise comparison of annotations ✓
- Which pairs do we compare?

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An alignment defines, which annotation of annotator 1 corresponds to which annotation of annotator 2 (if any)

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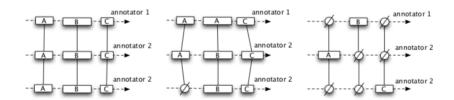


Figure: Different alignments between three annotators

Alignment: Two more ingredients

Calculate disagreement over a set of aligned individual annotations: Average

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Calculate disagreement over a set of aligned individual annotations: Average

$$\hat{\delta}(\hat{a}) = \frac{1}{|\hat{a}|} \sum_{(u,v) \in \hat{a}^2} d_{combi}(u,v)$$

with \hat{a} being a set of aligned annotations

Alignment: Two more ingredients

Calculate disagreement over a set of aligned individual annotations: Average

$$\hat{\delta}(\hat{a}) = \frac{1}{|\hat{a}|} \sum_{(u,v) \in \hat{a}^2} d_{combi}(u,v)$$

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with \hat{a} being a set of aligned annotations

► Calculate disagreement over a set of annotators: Average

$$\bar{\delta}(A) = \frac{1}{|x|} \sum_{i=1}^{|\hat{a}|} \hat{\delta}(\hat{a}_i)$$

with A being a set of annotators, and |x| the mean number of annotations per annotator

Alignment: Two more ingredients

Calculate disagreement over a set of aligned individual annotations: Average

$$\hat{\delta}(\hat{a}) = \frac{1}{|\hat{a}|} \sum_{(u,v) \in \hat{a}^2} d_{combi}(u,v)$$

with \hat{a} being a set of aligned annotations

Calculate disagreement over a set of annotators: Average

$$\bar{\delta}(A) = \frac{1}{|x|} \sum_{i=1}^{|\hat{a}|} \hat{\delta}(\hat{a}_i)$$

with A being a set of annotators, and |x| the mean number of annotations per annotator

• Alignment is created such that $\bar{\delta}(A)$ is minimal

Summary

- Gamma combines alignment and agreement calculation
- Core: Compare annotations pairwise, w.r.t.
 - their position
 - their categories
- Settable parameters
 - Dissimilarity of categories
 - Weighting between dissimilarity types
 - Position metric (SANTA: token numbers)
- Computationally expensive
- Implementation by Mathet et al. (2015) using ILP https://gamma.greyc.fr

Results

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Results

Note

The report shows Observed and Expected Disorder, i.e., the lower the better. Gamma scores represent agreement, i.e., the higher the better

All Annotations

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Results

Observed Disorder (all annotations)

SANTA_	1	2	3	4	5	6	7	8
Buechner	2.07	1.87	2.47	2.57	1.96	2.18	2.36	1.39
Chekhov	1.85	1.90	2.62	2.67	2.41	3.04	0.88	2.50
Henry	1.93	2.57	2.27	2.17	2.60	2.16	1.35	1.70
Kafka	1.38	1.03	2.68	1.50	1.49	1.82	1.27	1.00
Kleist	1.99	1.40	2.79	1.63	2.11	1.53	1.36	0.75
Lagerloef	1.81	2.75	2.36	2.45	1.50	2.81	0.91	0.67
Storm	2.07	1.62	2.48	2.35	2.12	1.97	1.42	1.45
Tieck	1.78	1.99	2.18	2.79	2.32	2.23	1.64	1.85
Min	1.38	1.03	2.18	1.50	1.49	1.53	0.88	0.67
Mean	1.86	1.89	2.48	2.27	2.06	2.22	1.40	1.41
Max	2.07	2.75	2.79	2.79	2.60	3.04	2.36	2.50
Stddev.	0.22	0.57	0.21	0.48	0.40	0.49	0.46	0.61

Results

Expected Disorder (all annotations)

SANTA_	1	2	3	4	5	6	7	8
Buechner	2.28	2.41	2.66	2.34	2.27	2.34	2.33	
Chekhov	2.10	2.24	2.47	2.54	2.51	2.88	1.96	
Henry	2.05	2.56	2.22	2.49	2.41	2.03	2.15	
Kafka	2.14				2.40	2.79	2.03	
Kleist	2.42	2.37	2.13			2.61	1.52	1.46
Lagerloef	1.82	2.82	2.38	2.26	2.03	2.39	1.63	
Storm	2.14	2.06	2.49	2.23	2.29	2.05	2.16	2.13
Tieck	2.02	2.05	2.55	2.25	2.39	2.08	2.38	
Min	1.82	2.05	2.13	2.23	2.03	2.03	1.52	1.46
Mean	2.12	2.36	2.41	2.35	2.33	2.40	2.02	1.80
Max	2.42	2.82	2.66	2.54	2.51	2.88	2.38	2.13
Stddev.	0.18	0.28	0.19	0.13	0.15	0.34	0.31	0.47

Resu

Inter-Annotator Agreement Gamma (all annotations)

SANTA_	1	2	3	4	5	6	7	8
Buechner	0.09	0.22	0.07	-0.10	0.14	0.07	-0.01	
Chekhov	0.12	0.15	-0.06	-0.05	0.04	-0.05	0.55	
Henry	0.06	0.00	-0.02	0.13	-0.08	-0.06	0.37	
Kafka	0.36				0.38	0.35	0.37	
Kleist	0.18	0.41	-0.31			0.41	0.11	0.49
Lagerloef	0.01	0.02	0.01	-0.08	0.26	-0.18	0.44	
Storm	0.03	0.21	0.00	-0.05	0.07	0.04	0.34	0.32
Tieck	0.12	0.03	0.15	-0.24	0.03	-0.07	0.31	
Min	0.01	0.00	-0.31	-0.24	-0.08	-0.18	-0.01	0.32
Mean	0.12	0.15	-0.02	-0.07	0.12	0.06	0.31	0.40
Max	0.36	0.41	0.15	0.13	0.38	0.41	0.55	0.49
Stddev.	0.11	0.15	0.14	0.12	0.15	0.21	0.18	0.12

Table: Inter-Annotator Agreement Gamma (the higher the better)

Results

Own vs. Foreign Annotations

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Observed Disorder (own vs. foreign)

SANTA_	1	2	3	4	5	6	7	8
Buechner	1.72	1.64	2.00	1.99	1.47	1.39	1.49	1.11
Chekhov	1.54	1.33	1.90	1.96	1.68	2.00	0.56	1.84
Henry	1.61	1.64	2.00	1.42	1.00	1.26	1.20	2.12
Kafka	1.25	1.07	1.62	1.50	1.22	1.18	0.80	0.00
Kleist	1.83	0.91	1.95	1.37	2.00	1.44	0.00	0.00
Lagerloef	1.75	1.73	2.05	1.82	1.85	2.50	0.50	1.00
Storm	1.94	1.17	1.92	1.89	1.55	1.68	1.23	1.59
Tieck	1.61	1.55	1.94	2.00	1.75	1.85	1.80	0.93
Min	1.25	0.91	1.62	1.37	1.00	1.18	0.00	0.00
Mean	1.65	1.38	1.92	1.74	1.57	1.66	0.95	1.07
Max	1.94	1.73	2.05	2.00	2.00	2.50	1.80	2.12
Stddev.	0.22	0.57	0.21	0.48	0.40	0.49	0.46	0.61

Expected Disorder (own vs. foreign)

SANTA_	1	2	3	4	5	6	7	8
Buechner	1.68	1.81	1.60	1.79	1.83	1.44	1.49	1.55
Chekhov	1.66	1.67	1.79	1.56	1.81	1.51	1.50	1.70
Henry	1.71	1.78	1.59	1.83	1.55	1.29	1.66	1.67
Kafka	1.61	1.80	1.50	1.56	1.91	1.70	1.42	
Kleist	1.66	1.65	1.39	1.56		1.81	0.00	0.00
Lagerloef	1.58	1.90	1.51	1.52	1.68	1.93	1.32	
Storm	1.59	1.64	1.65	1.69	1.62	1.63	1.64	1.63
Tieck	1.59	1.60	1.82	1.70	1.81	1.58	1.81	1.38
Min	1.58	1.60	1.39	1.52	1.55	1.29	0.00	0.00
Mean	1.64	1.73	1.61	1.65	1.75	1.61	1.35	1.32
Max	1.71	1.90	1.82	1.83	1.91	1.93	1.81	1.70
Stddev.	0.18	0.28	0.19	0.13	0.15	0.34	0.31	0.47

Results

Inter-Annotator Agreement Gamma (own vs. foreign)

SANTA_	1	2	3	4	5	6	7	8
Buechner	-0.02	0.09	-0.25	-0.11	0.20	0.04	0.00	0.28
Chekhov	0.07	0.21	-0.06	-0.26	0.07	-0.32	0.63	-0.08
Henry	0.06	0.08	-0.26	0.22	0.36	0.02	0.28	-0.27
Kafka	0.23	0.41	-0.08	0.04	0.36	0.31	0.44	1.00
Kleist	-0.10	0.45	-0.40	0.12		0.20	1.00	1.00
Lagerloef	-0.10	0.09	-0.36	-0.19	-0.10	-0.30	0.62	
Storm	-0.21	0.29	-0.17	-0.11	0.05	-0.03	0.25	0.02
Tieck	-0.01	0.03	-0.06	-0.18	0.03	-0.17	0.00	0.32
Min	-0.21	0.03	-0.40	-0.26	-0.10	-0.32	0.00	-0.27
Mean	-0.01	0.21	-0.21	-0.06	0.14	-0.03	0.40	0.33
Max	0.23	0.45	-0.06	0.22	0.36	0.31	1.00	1.00
Stddev.	0.11	0.15	0.14	0.12	0.15	0.21	0.18	0.12

Table: Inter-Annotator Agreement Gamma (the higher the better)

Own vs. Student Annotations

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Observed Disorder (own vs. student)

SANTA_	1	2	3	4	5	6	7	8
Buechner	1.48	1.07	1.77	1.00	1.53	1.99	1.87	1.33
Chekhov	1.70	1.69	1.86	1.33	1.72	2.50	0.84	1.33
Henry	1.59	2.11	1.59	1.67	1.69	1.75	0.97	1.50
Kafka	1.33	1.00	2.00	0.00	1.16	2.25	0.90	1.00
Kleist	1.62	1.31	1.95	1.00	1.73	1.25	1.36	0.75
Lagerloef	1.58	2.00	1.59	0.67	0.63	2.01	0.95	0.00
Storm	1.47	1.30	1.82	0.57	1.70	1.71	1.05	0.42
Tieck	1.52	1.49	1.54	1.32	1.90	1.70	0.92	1.00
Min	1.33	1.00	1.54	0.00	0.63	1.25	0.84	0.00
Mean	1.53	1.50	1.76	0.94	1.51	1.90	1.11	0.92
Max	1.70	2.11	2.00	1.67	1.90	2.50	1.87	1.50
Stddev.	0.22	0.57	0.21	0.48	0.40	0.49	0.46	0.61

Results

Expected Disorder (own vs. student)

SANTA_	1	2	3	4	5	6	7	8
Buechner	1.58	1.75	1.86		1.65	1.75	1.64	1.35
Chekhov	1.66	1.72	1.67	1.34	1.76	1.88	1.57	1.30
Henry	1.51	1.91	1.63	1.77	1.48	1.70	1.60	1.44
Kafka	1.77	1.80	1.64		1.68	2.09	1.54	
Kleist	1.70	1.74	1.61	1.43	1.78	1.82	1.52	1.51
Lagerloef	1.40	1.94	1.68	1.19	1.43	1.41	1.37	1.50
Storm	1.49	1.48	1.77	1.50	1.67	1.70	1.66	1.55
Tieck	1.49	1.45	1.88	1.36	1.74	1.64	1.71	1.44
Min	1.40	1.45	1.61	1.19	1.43	1.41	1.37	1.30
Mean	1.58	1.72	1.72	1.43	1.65	1.75	1.58	1.44
Max	1.77	1.94	1.88	1.77	1.78	2.09	1.71	1.55
Stddev.	0.18	0.28	0.19	0.13	0.15	0.34	0.31	0.47

Results

Inter-Annotator Agreement Gamma (own vs. student)

SANTA_	1	2	3	4	5	6	7	8
Buechne	r 0.07	0.39	0.05		0.07	-0.14	-0.14	0.01
Chekhov	-0.02	0.02	-0.11	0.01	0.03	-0.33	0.47	-0.02
Henry	-0.05	-0.10	0.02	0.06	-0.14	-0.03	0.39	-0.04
Kafka	0.25	0.44	-0.22	1.00	0.31	-0.07	0.42	
Kleist	0.05	0.25	-0.21	0.30	0.02	0.32	0.11	0.50
Lagerloe	f-0.13	-0.03	0.06	0.44	0.56	-0.43	0.31	1.00
Storm	0.01	0.12	-0.03	0.62	-0.02	-0.01	0.37	0.73
Tieck	-0.02	-0.03	0.18	0.03	-0.09	-0.04	0.47	0.30
Min	-0.13	-0.10	-0.22	0.01	-0.14	-0.43	-0.14	-0.04
Mean	0.02	0.13	-0.03	0.35	0.09	-0.09	0.30	0.35
Max	0.25	0.44	0.18	1.00	0.56	0.32	0.47	1.00
Stddev.	0.11	0.15	0.14	0.12	0.15	0.21	0.18	0.12

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SANTA_	1	2	3	4	5	6	7	8
Buechner	1.67	1.67	1.97	1.86	1.68	1.88	1.52	1.02
Chekhov	1.38	1.69	1.61	1.75	2.17	2.07	0.88	1.51
Henry	1.52	2.17	2.00	1.77	1.83	1.66	1.20	1.52
Kafka	1.17	1.08	2.00	1.50	1.41	2.33	1.18	1.00
Kleist	1.48	0.91	1.50	1.25	1.68	1.21	0.00	0.00
Lagerloef	1.06	1.96	2.00	1.79	1.73	1.84	0.84	1.00
Storm	1.89	1.56	2.02	1.93	1.80	1.50	1.31	1.38
Tieck	1.41	1.91	2.00	2.00	1.65	1.73	1.83	1.20
Min	1.06	0.91	1.50	1.25	1.41	1.21	0.00	0.00
Mean	1.45	1.62	1.89	1.73	1.74	1.78	1.09	1.08
Max	1.89	2.17	2.02	2.00	2.17	2.33	1.83	1.52
Stddev.	0.22	0.57	0.21	0.48	0.40	0.49	0.46	0.61

Results

Expected Disorder (foreign vs. student)

SANTA_	1	2	3	4	5	6	7	8
Buechner	1.61	1.82	1.52	1.43	1.76	1.85	1.51	1.20
Chekhov	1.57	1.68	1.67	1.77	1.89	1.88	1.55	
Henry	1.55	2.06	1.55	1.72	1.60	1.49	1.70	1.52
Kafka	1.57	2.43		1.61	1.78	2.17	1.65	
Kleist	1.69	1.65		1.39		1.78	0.00	0.00
Lagerloef	1.14	1.80	1.46	1.53	1.61	1.81	1.44	
Storm	1.63	1.74	1.65	1.65	1.71	1.50	1.69	1.71
Tieck	1.63	1.73	1.86	1.51	1.84	1.55	1.79	1.49
Min	1.14	1.65	1.46	1.39	1.60	1.49	0.00	0.00
Mean	1.55	1.86	1.62	1.58	1.74	1.75	1.42	1.19
Max	1.69	2.43	1.86	1.77	1.89	2.17	1.79	1.71
Stddev.	0.18	0.28	0.19	0.13	0.15	0.34	0.31	0.47

Results

Inter-Annotator Agreement Gamma (foreign vs. student)

SANTA_	1	2	3	4	5	6	7	8
Buechne	r-0.04	0.09	-0.29	-0.30	0.04	-0.01	-0.01	0.15
Chekhov	0.12	0.00	0.04	0.01	-0.15	-0.10	0.44	
Henry	0.02	-0.05	-0.29	-0.03	-0.15	-0.11	0.29	0.01
Kafka	0.25	0.56		0.07	0.21	-0.08	0.29	
Kleist	0.13	0.45		0.10		0.32	1.00	1.00
Lagerloet	f 0.06	-0.09	-0.37	-0.17	-0.08	-0.02	0.42	
Storm	-0.16	0.10	-0.23	-0.17	-0.05	0.00	0.23	0.19
Tieck	0.13	-0.10	-0.08	-0.32	0.10	-0.12	-0.02	0.20
Min	-0.16	-0.10	-0.37	-0.32	-0.15	-0.12	-0.02	0.01
Mean	0.07	0.12	-0.20	-0.10	-0.01	-0.01	0.33	0.31
Max	0.25	0.56	0.04	0.10	0.21	0.32	1.00	1.00
Stddev.	0.11	0.15	0.14	0.12	0.15	0.21	0.18	0.12

Table: Inter-Annotator Agreement Gamma (the higher the better)

Comments

- Obvious ways to boost scores
 - Simple schemes without many features get higher scores (in addition to being easier to annotate)
- Some guidelines include more narrative phenomena than 'levels'
 - Can also lead to lower IAA