

Three of a kind?

Multi-alignment of amino-acids, sounds and words

Michael Cysouw
Philipps-Universität Marburg

Multi-alignment

- A. Biological Phylogenetics
(alignment of nucleotides/amino-acids)
- B. Historical Linguistics
(alignment of sounds in words)
- C. Functional Language Comparison
(alignment of words in sentences)

A: Biological Phylogenetics

- Multi-alignment methods actively developed since the 1980s
- Start with pairwise alignment, then build up multiple alignment starting from most similar pairs (using “guide tree”)

Multi-alignment of nucleotides (4-letter alphabet)

AliView - Woodsia_chloroplast_min4_20131109_v2.excluded.nexus

File Edit Selection View Align Primer External commands Help

Color palette: A A

	130	1140	1150	1160	1170	1180	1190
Thelypteris_palustris_2x_F14	c	g	a	a	t	t	t
Woodsia_alpina_4x_F22_F89_F95_F1	c	g	a	a	t	t	t
Woodsia_alpina_4x_F4_F135_F137	c	g	a	a	t	t	t
Woodsia_andersonii_8x_F56_F115_F	c	g	a	a	t	t	t
Woodsia_andersonii_8x_F75	c	g	a	a	t	t	t
Woodsia_canescens_2x4x_F66	c	g	a	a	t	t	t
Woodsia_cochisensis_4x_F48_F152_I	c	g	a	a	t	t	t
Woodsia_cycloloba_8x_F72	c	g	a	a	t	t	t
Woodsia_elongata_2x4x_F1_F92	c	g	a	a	t	t	t
Woodsia_fragilis_4x_F19_F114_F227	c	g	a	a	t	t	t
Woodsia_gabella_2x_F86	c	g	a	a	t	t	t
Woodsia_gabella_2x_F88_F167	c	g	a	a	t	t	t
Woodsia_gracilima_2x_F55	c	g	a	a	t	t	t
Woodsia_ilvensis_2x_F3_F21_F77_F7	c	g	a	a	t	t	t
Woodsia_indusiosa_4x_F91_F169	c	g	a	a	t	t	t
Woodsia_intermedia_4x_F71	c	g	a	a	t	t	t
Woodsia_lanosa_8x_F119	c	g	a	a	t	t	t
Woodsia_macrochlaena_2x_F6	c	g	a	a	t	t	t
Woodsia_manchuriensis_2x4x_F47_F	c	g	a	a	t	t	t
Woodsia_mollis_3x4x_F18_F143_F25	c	g	a	a	t	t	t
Woodsia_burgessiana_syn_montevide	c	g	a	a	t	t	t
Woodsia_montevidensis_ecuador_2x	c	g	a	a	t	t	t
Woodsia_neomexicana_4x_F68_F154	c	g	a	a	t	t	t
Woodsia_obtusa_2x4x_F70	c	g	a	a	t	t	t
Woodsia_okamotoi_4x8x_F176	c	g	a	a	t	t	t
Woodsia_oregana_ssp_oregana_2x_I	c	g	a	a	t	t	t
Woodsia_aff_phillipsii_4x_F69_F161	c	g	a	a	t	t	t
Woodsia_plummerae_4x_F31_F44_F	c	g	a	a	t	t	t
Woodsia_polystichoides_2x_F11	c	g	a	a	t	t	t
Woodsia_rosthorniana_4x_F54_F94	c	g	a	a	t	t	t
Woodsia_scopulina_ssp_scopulina_2	c	g	a	a	t	t	t

Multi-alignment of amino-acids (20-letter alphabet)

		*	:	.	*	:	:	.
Q5E940_BOVIN	-----MPREDRATWKSNYFLKIIQLLDDYPKCFIVGADNVGSKQMQQIRMSLRGK-AVVLMGKNTMMRKAIRGHLENN--PALE							
RLAO_HUMAN	-----MPREDRATWKSNYFLKIIQLLDDYPKCFIVGADNVGSKQMQQIRMSLRGK-AVVLMGKNTMMRKAIRGHLENN--PALE							
RLAO_MOUSE	-----MPREDRATWKSNYFLKIIQLLDDYPKCFIVGADNVGSKQMQQIRMSLRGK-AVVLMGKNTMMRKAIRGHLENN--PALE							
RLAO_RAT	-----MPREDRATWKSNYFLKIIQLLDDYPKCFIVGADNVGSKQMQQIRMSLRGK-AVVLMGKNTMMRKAIRGHLENN--PALE							
RLAO_CHICK	-----MPREDRATWKSNYFMKIIQLLDDYPKCFVVGADNVGSKQMQQIRMSLRGK-AVVLMGKNTMMRKAIRGHLENN--PALE							
RLAO_RANSY	-----MPREDRATWKSNYFLKIIQLLDDYPKCFIVGADNVGSKQMQQIRMSLRGK-AVVLMGKNTMMRKAIRGHLENN--SALE							
Q7ZUG3_BRARE	-----MPREDRATWKSNYFLKIIQLLDDYPKCFIVGADNVGSKQMQTIRLSLRGK-AVVLMGKNTMMRKAIRGHLENN--PALE							
RLAO_ICTPU	-----MPREDRATWKSNYFLKIIQLLNDYPKCFIVGADNVGSKQMQTIRLSLRGK-AIVLMGKNTMMRKAIRGHLENN--PALE							
RLAO_DROME	-----MRENKAQYFPIKVVELFDEFPKCFIVGADNVGSKQMQNIRTSLRGL-AVVLMGKNTMMRKAIRGHLENN--PQLE							
RLAO_DICDI	-----MSGAG-SKRKKLFIEKATKLFTTYDKMIVAEADFVGSSQLQKIRKSIRGI-GAVLMGKTMIRKVIRDLADSK--PELD							
Q54LP0_DICDI	-----MSGAG-SKRKNVFIEKATKLFTTYDKMIVAEADFVGSSQLQKIRKSIRGI-GAVLMGKTMIRKVIRDLADSK--PELD							
RLAO_PLAF8	-----MAKLSQQKKQMYIEKLSSLIQQYSKILIVHVDNVGSNQMASVRKSLRGK-ATILMGKNTIRRTALKKNLQAV--PQIE							
RLAO_SULAC	-----MIGLAVTTKKIAKWKVDEVAEELTEKLKTHKTIIIANIEGFPADKLHEIRKKLRGK-ADIKVTKNLFNIALKNAG---YDTK							
RLAO_SULTO	-----MRIMAVITQERKIAKWKIEEVKELEQKLREYHTIIIANIEGFPADKLHDIRKKMRGM-AEIKVTKNTLFGIAAKNAG---LDVS							
RLAO_SULSO	-----MKRLALALKQRKVASWKEEVKELTELIKNSNTILIGNLEGFPADKLHEIRKKLRGK-ATIKVTKNTLFKIAAKNAG---IDIE							
RLAO_AERPE	MSVVSLEVQMYKREKPIPEWKTLMRLELEELFSKHRVVLFAADLTGPTFVVVQRVKKLWKK-YPMMVAKRIILRAMKAAGLE--LDDN							
RLAO_PYRAE	-MMLAIGKRRYVRTRQYPARKVKIVSEATELLQKYPYVFLFDLHGLSSRILHEYRYRLRRY-GVIKIIKPTLFKIAFTKVYGG--IPAE							
RLAO_METAC	-----MAEERHHTEHIPQWKKDEIENIKELIQSHKVFGMVRIEGILATKMQKIRRDLKDV-AVLKVSRNTLTERALNQLG---ETIP							
RLAO_METMA	-----MAEERHHTEHIPQWKKDEIENIKELIQSHKVFGMVRIEGILATKIQKIRRDLKDV-AVLKVSRNTLTERALNQLG---ESIP							
RLAO_ARCFU	-----MAAVRGS--PPEYKVRAVEEIKRMISSSKPVVAIVSFRNVPAGQMOKIRREFRGK-AEIKVVKNTLLERALDALG---GDYL							
RLAO_METKA	MAVKAKGQPPSGYEPKVAEWKRREVKELKELMDEYENVGLVDLEGIPAPQLQEIRAKLRRERDTIIRMSRNTLMRIAEEKLDER--PELE							
RLAO_METTH	-----MAHVAEWKKKEVQELHDLIKGYEVVGIANLADIPARQLQKMRQTLRDS-ALIRMSKKTLISLALEKAGREL--ENVD							
RLAO_METTL	-----MITAESEHKIAPWKIEEVNKLKELLKNGQIVALVDMMEVPARQLQEIRDKIR-GTMTLKMSRNTLIERAIKEVAEETGNPEFA							
RLAO_METVA	-----MIDAKSEHKIAPWKIEEVNALKELLKSANVIALIDMMEVPAVQLQEIRDKIR-DQMTLKMSRNTLIKRAVEEEVAEETGNPEFA							
RLAO_METJA	-----METKVKAHVAPWKIEEVKTLKGLIKSKPVVAIVDMMMDVPAQQLQEIRDKIR-DKVKLRLMSRNTLIIRALKEAAEELNNPKLA							
RLAO_PYRAB	-----MAHVAEWKKKEVEELANLIKSYPPVIALVDVSSMPAYPLSQMRRLIRENGLLRVSRNTLIELAIKAAQELGKPELE							
RLAO_PYRHO	-----MAHVAEWKKKEVEELAKLIKSYPPVIALVDVSSMPAYPLSQMRRLIRENGLLRVSRNTLIELAIKAAKELGKPELE							
RLAO_PYRFU	-----MAHVAEWKKKEVEELANLIKSYPPVVALVDVSSMPAYPLSQMRRLIRENNGLLRVSRNTLIELAIKVAQELGKPELE							
RLAO_PYRKO	-----MAHVAEWKKKEVEELANIKSYPVIALVDVAGVPAYPLSKMRDKLR-GKALLRVSNTLIELAIKRAAQELGQPELE							
RLAO_HALMA	-----MSAESERKTETIPEWKQEEVDAIEMIESYESVGVNVIAGIPSRLQLQDMRRDLHGT-AELRVSRNTLLERALDDVD---DGLE							
RLAO_HALVO	-----MSESEVRQTEVIPQWKREEVDELDFIESYESVGVVGVAGIPSRLQLQSMRREIHGS-AAVRMSRNTLVNRALDEVN---DGFE							
RLAO_HALSA	-----MSAEEQRTTEEVPFWKRQEVAELVDSLLETYDSVGVVNTGIPSKQLQDMRRGLHGQ-AALRMSRNTLLVRALEEAG---DGLD							
RLAO_THEAC	-----MKEVSQQKKELVNEITQRIKASRSVAIVDTAGIRTROIQDIRGKNGK-INLKVIKKTLLFKALENLGD---EKLS							
RLAO_THEVO	-----MRKINPKKEIVSELAQDITSKAVAAIVDIKGVRTRQMQDIRAKNRDK-VKIKVVVKTLLFKALDSIND---EKLT							
RLAO_PICTO	-----MTEPAQWKIDFVKNLENEINSRKVAAIVSIKGLRNNEFQKIRNSIRDK-ARIKVSRARLLRAIENTGK---NNIV							

B: Historical Linguistics

- Multi-alignment is just a fancy name for sound correspondences
- Each sound correspondence is “aligned” in a column, possibly adding empty cells
- It is a useful and consistent way to represent comparative data (both between languages or dialects)
- First automatic multi-alignments by Kondrak (2009) and Prokić (2009). Currently furthest developed by Mattis List (LingPy)

Language	IDS	meaning	alignment						
Pilagá	15.810/15.820	heavy/light	d	e	s	a	l	i	
Toba	15.810	heavy	d	e	s	a	l ^y	i	
Mocoví	15.810/15.820	heavy/light	r	e	s	a	l ^y	i	
Pilagá	9.440	build	n	?	o	v	o	–	s e g e m
Toba	9.440	build	n	?	o	v	o	o	š i g e m
Mocoví	9.440	build	n	o	?	v	o	n	š i g i m

Affe (German Dialect data)

LOCATION	WORD
Aachen	a:ph
Adorf	q:b ^h ə
Ahrbergen	o→çphə
Albersloh	a:p ^h ə
Allna	aɸh
Altenberg	Λfɛ
Altentrüdin	af
Altlandsberg	a'fə'
Altwarp	o:ph
Astfeld	v ^c :p ^h ə
Atzendorf	afɛ
Ballhausen	Λ'fə
Bardenfleth	ɔ:pΦ
Barssel	ä:p ^h ə
Bempflingen	af:
Bennin	ɔp ^h
Billingsbach	af
Bockelwitz	Λvə
Bonn	a:p'
Borstendorf	χf:
Breddin	v:ph
Brelingen	qfβə
Bremscheid	v ^c :phə
...	...

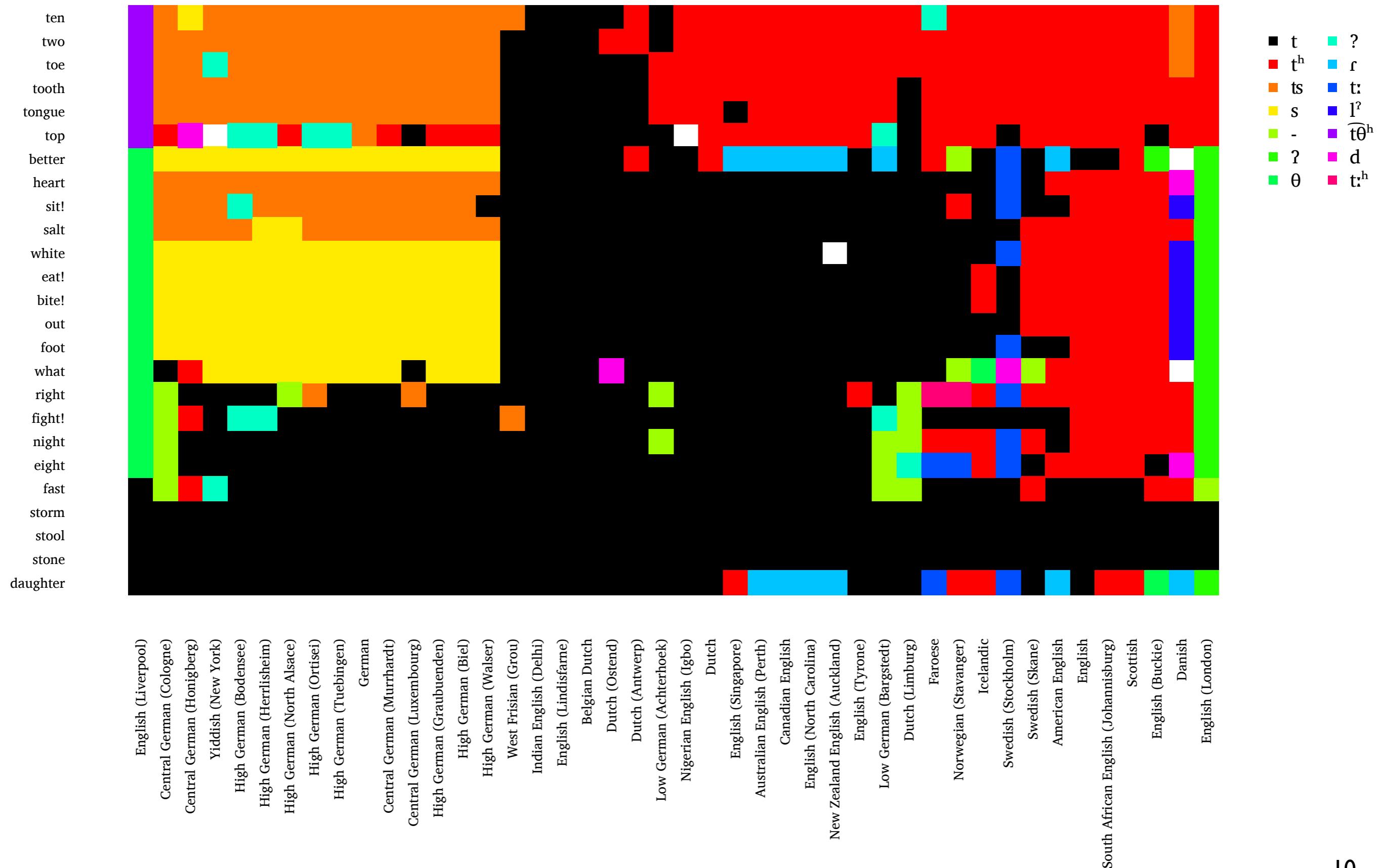
A	FF	E
a:	ph	-
ä:	b ^h	ə
o→ç	ph	ə
a:	p ^h	ə
a	ɸh	-
Λ	f	ɛ
a	f	-
a'	f	ə'
o:	ph	-
v ^c :	p ^h	ə
a	f	ɛ
Λ'	f	ə
ɔ:	pΦ	-
ä:	p ^h	ə
a	f:	-
ɔ	p ^h	-
a	f	-
Λ	v	ə
a:	p'	-
χ	f:	-
v:	ph	-
ä	fβ	ə
v ^c :	ph	ə
...

tongue

American English	t ^h	ʌ	ə	-	-
Canadian English	t ^h	ʌ	ə	-	-
Central German (Cologne)	ts	ʊ	ə	-	-
Central German (Honigberg)	ts	aɪ	-	-	-
Central German (Luxembourg)	ts	ɔ̄	ə	-	-
Central German (Murrhardt)	ts	ʊ	ə	-	-
Danish	t ^h	ʊ	ə	-	ø
Dutch (Antwerp)	t	ʌ	ə	-	-
Belgian Dutch	t	ʊ	ə	-	-
Dutch (Limburg)	t	ʊ	ə	-	-
Dutch (Ostend)	t	ʊ	ə	-	ø
Dutch	t ^h	ʊ	ə	-	-
New Zealand English (Auckland)	t ^h	ə	ə	-	-
English (Buckie)	t ^h	ʊ	ə	-	-
Indian English (Delhi)	t	ə	ə	-	-
Nigerian English (Igbo)	t ^h	v	ə	g	-
South African English (Johannisburg)	t ^h	ɛ	ə	-	-
English (Lindisfarne)	t	ɔ̄	ə	-	-
English (Liverpool)	tθ ^h	ʊ	ə	g	-
English (London)	t ^h	ə	ə	-	-
English (North Carolina)	t ^h	ɪə	ə	-	-
Australian English (Perth)	t ^h	ə	ə	-	-
English (Singapore)	t	ə	ə	-	-
English	t ^h	ə	ə	-	-
English (Tyrone)	t ^h	ɔ̄	ə	-	-
Faroese	t ^h	ʊ	ə	k	a
German	ts	ʊ	ə	-	ø
High German (North Alsace)	ts	ʊ	ə	-	-
High German (Biel)	ts	ʊ	ə	-	ø
High German (Bodensee)	ts	ʊ	ə	-	ø
High German (Graubuenden)	ts	ʊ	ə	g	æ
High German (Herrlisheim)	ts	ʊ	ə	-	-
High German (Ortisei)	ts	ʊ	ə	g	ɛ
High German (Tuebingen)	ts	u	ə	g	-
High German (Walser)	ts	ʊ	ə	g	ø
Icelandic	t ^h	ʊ	ə	k	a
Low German (Achterhoek)	t ^h	ʊ	ə	-	ø
Low German (Bargstedt)	t ^h	ʊ	ə	-	-
Norwegian (Stavanger)	t ^h	ʊ	ə	-	ø
Scottish	t ^h	ʌ	ə	-	-
Swedish (Skane)	t ^h	øy	ə	j	æ
Swedish (Stockholm)	t ^h	ʊ	ə	-	ø
West Frisian (Grou)	t	ɔ̄	ə	-	ø
Yiddish (New York)	ts	u	ə	g	-

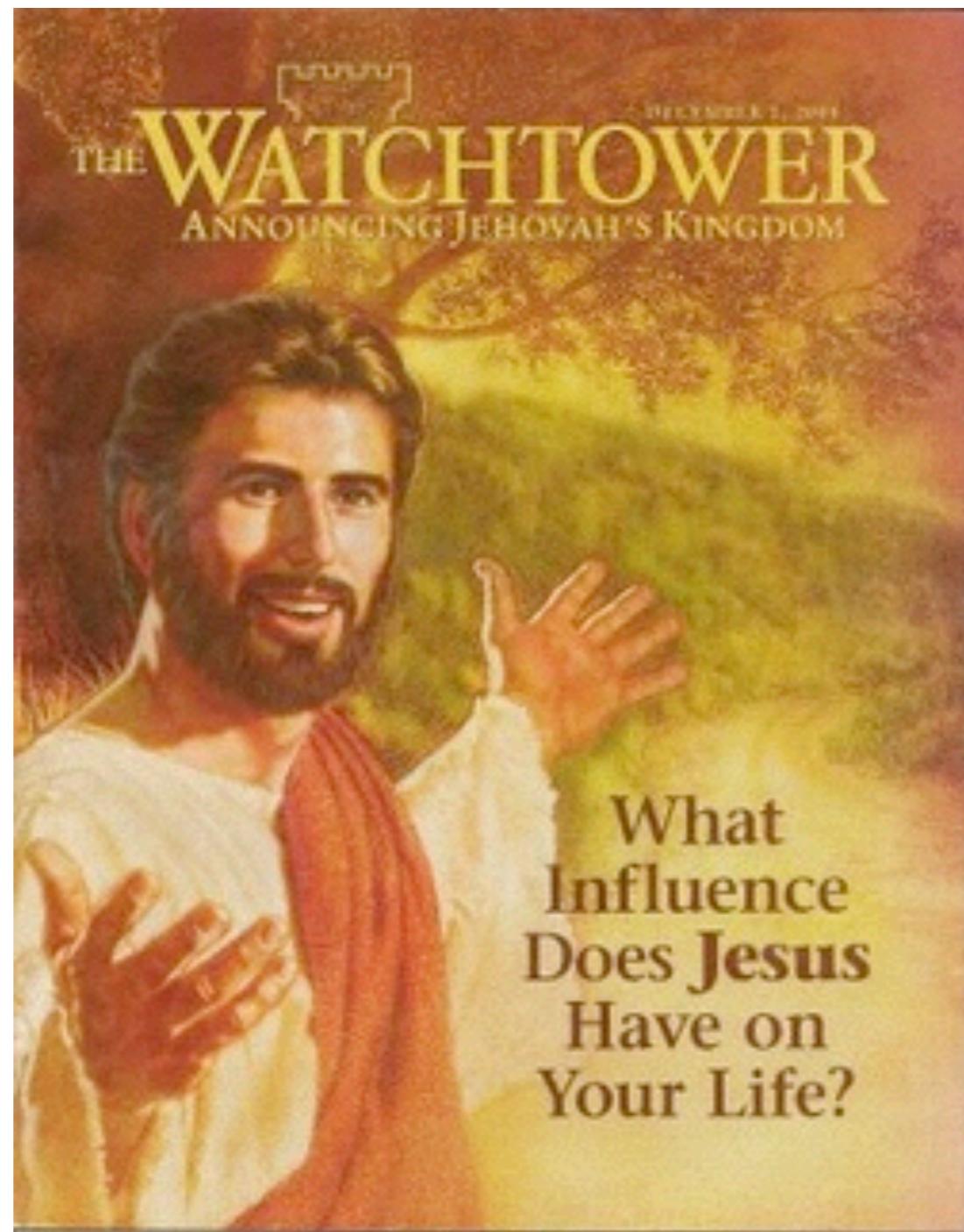
(Data from Paul Haggerty,
aligned by the LingPy library
maintained by Mattis List)

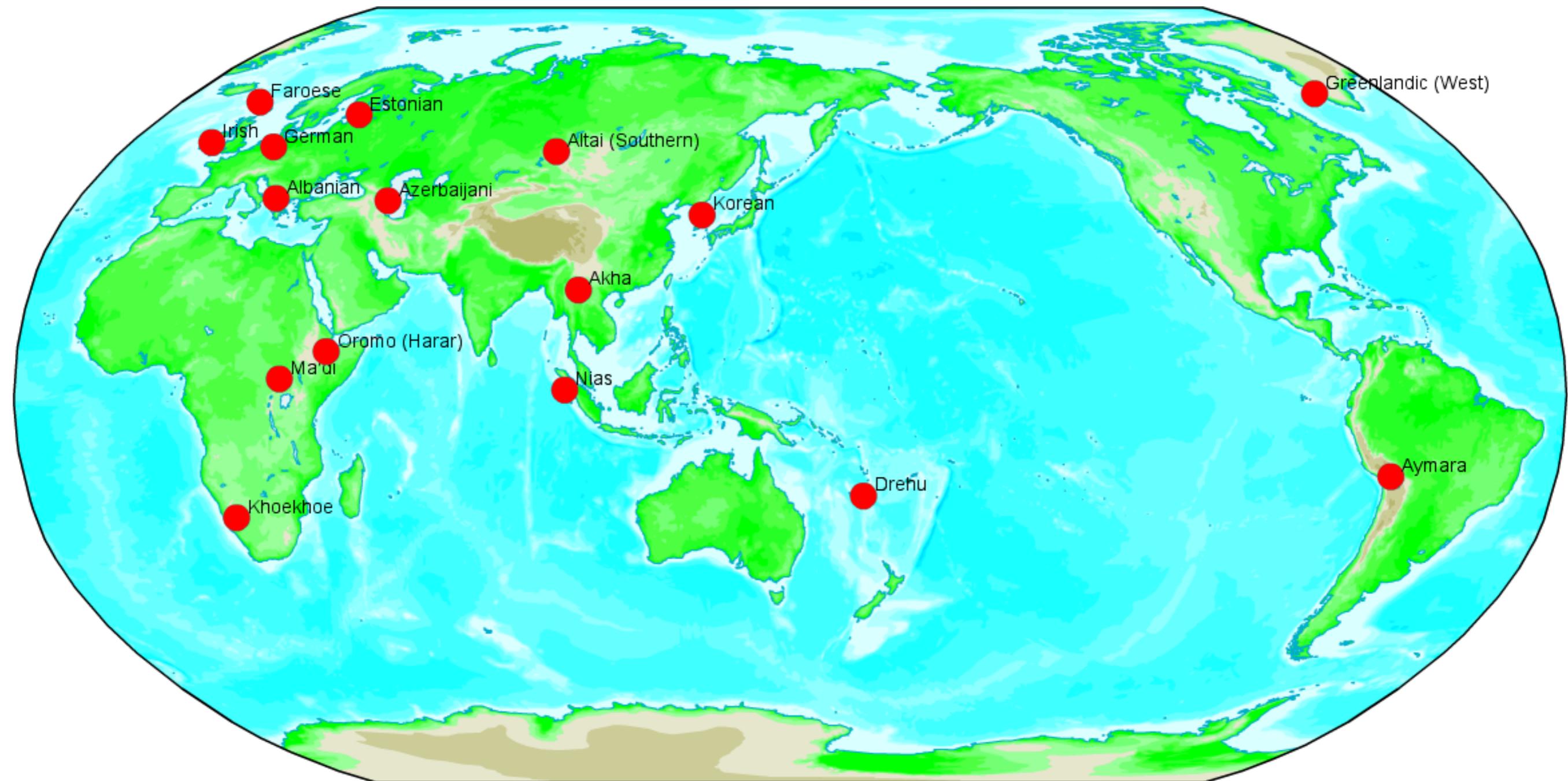
Correspondences with high frequency of [t]



C: Functional Language Comparison

- Comparing morphosyntactic patterns in languages all over the world
- “Language Typology”
- Traditional problem how to compare disparate languages.
- Solution: Multi-alignment



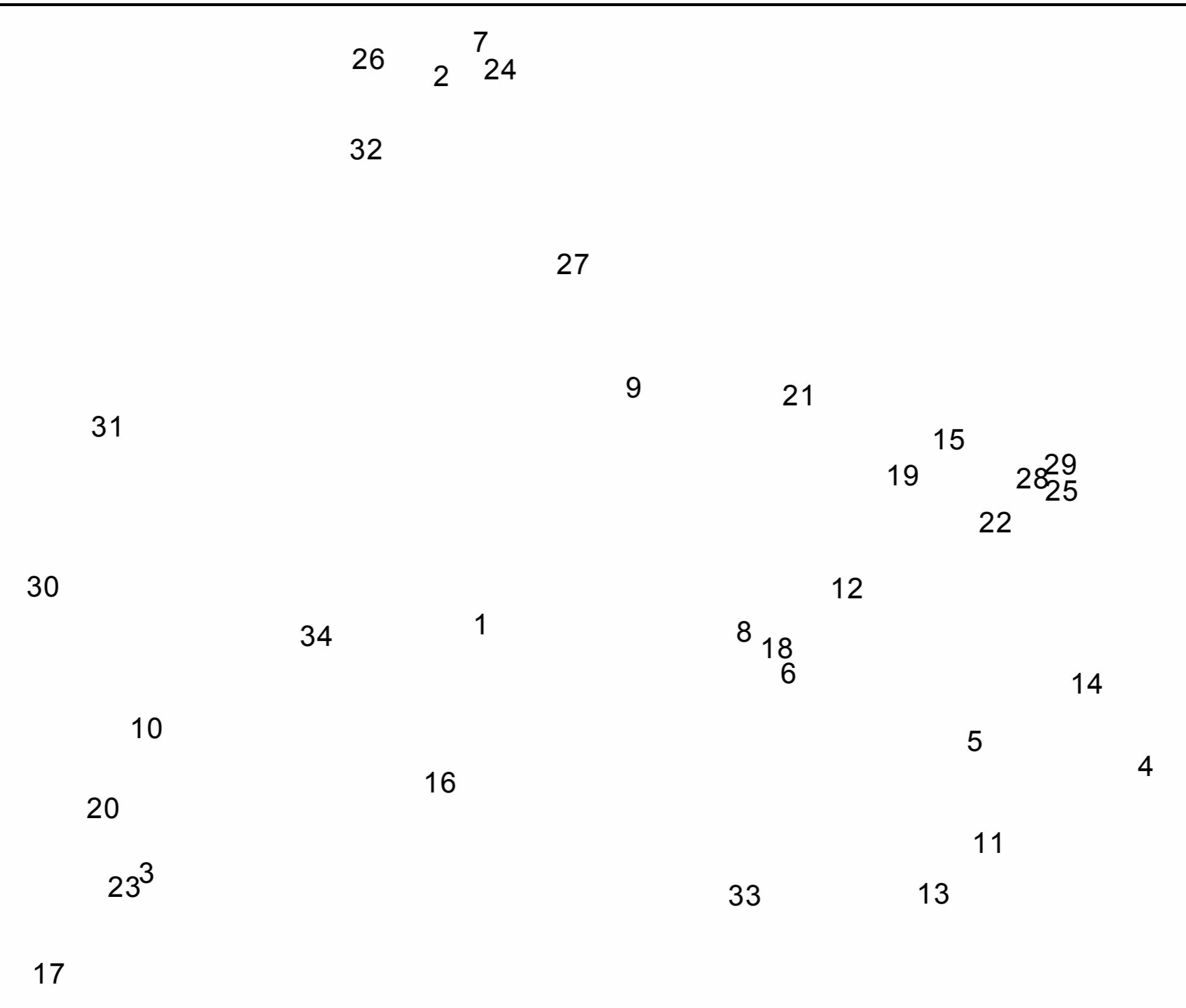


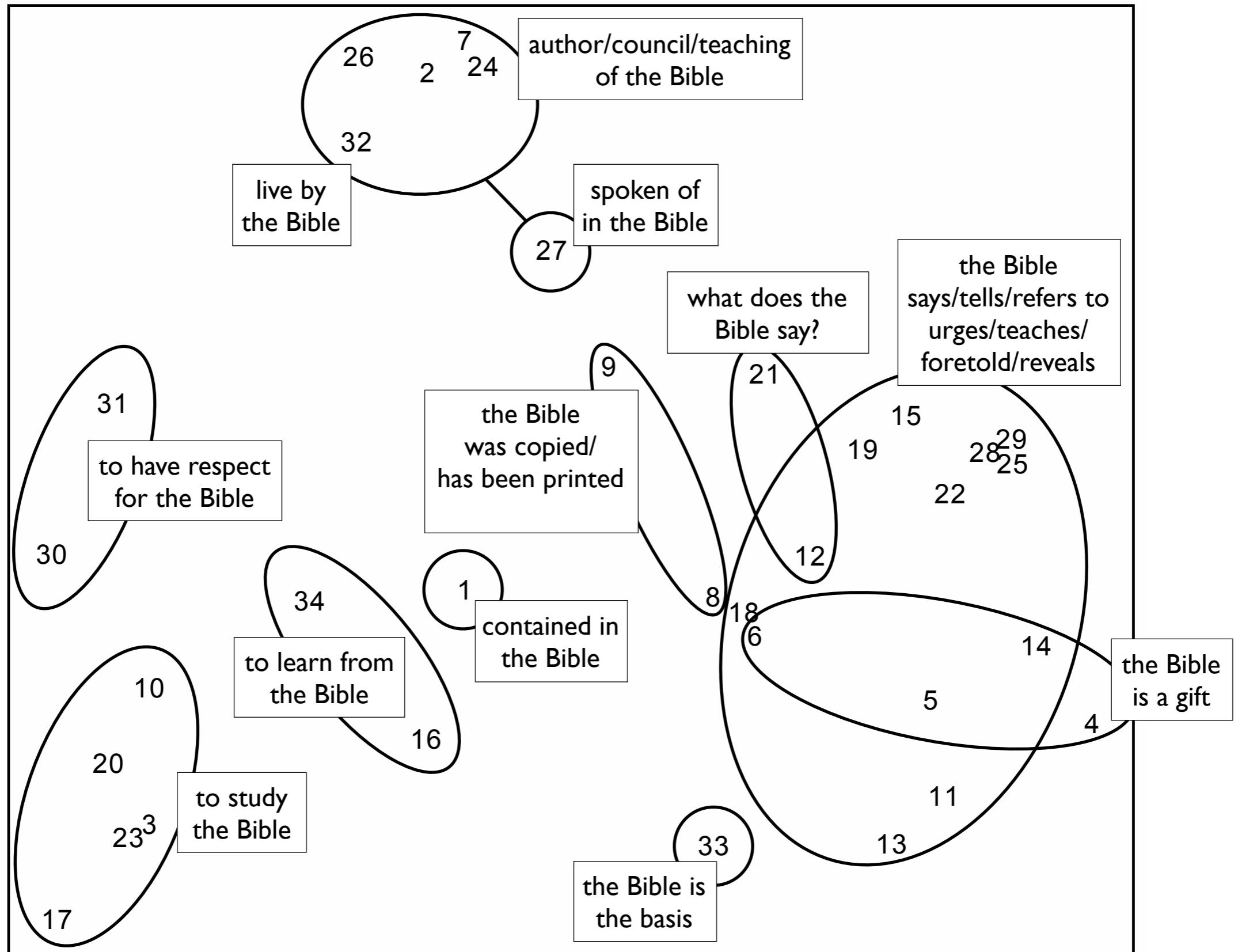
- 1 What important information is contained in the Bible?
- 2 Who is the Bible's author?
- 3 Why should you study the Bible?
- 4 The Bible is a precious gift from God.
- 5 The Bible alone tells us what we must do to please God.
- 6 The Bible was written by some 40 different men over a period of 1,600 years, beginning in 1513 B.C.E.
- 7 So God in heaven, not any human on earth, is the Author of the Bible.
- 8 God made sure that the Bible was accurately copied and preserved.
- 9 More Bibles have been printed than any other book.
- 10 Not everyone will be happy to see you studying the Bible, but do not let that stop you.
- 11 But the Bible tells us that there is only one TRUE God.
- 12 But when the Bible was written, the name Jehovah appeared in it some 7,000 times
- 13 God is a Spirit, says the Bible.
- 14 The Bible reveals Jehovah's personality to us.
- 15 The Bible tells us that he is also merciful, kind, forgiving, generous, and patient.
- 16 We learn about God from creation and from the Bible.
- 17 Another way we can learn about God is by studying the Bible.
- 18 By disobeying God's command, the first man, Adam, committed what the Bible calls sin.
- 19 This is what the Bible refers to as the ransom.
- 20 Some of your loved ones may become very angry because you are studying the Bible.
- 21 What is the Bible's view of separation and of divorce?
- 22 The Bible says that a husband is the head of his family.
- 23 Parents need to spend time with their children and study the Bible with them,
- 24 When marriage mates have problems getting along together, they should try to apply Bible counsel.
- 25 The Bible urges us to show love and to be forgiving.
- 26 But God does not approve of them if they come from false religion or are against Bible teachings.
- 27 The only two birthday celebrations spoken of in the Bible were held by persons who did not worship Jehovah.
- 28 The Bible teaches that only a few people are on the narrow road to life.
- 29 The Bible foretold that after the death of the apostles, ...
- 30 True Christians love one another, respect the Bible, and preach about God's Kingdom.
- 31 Another mark of true religion is that its members have a deep respect for the Bible.
- 32 They try to live by the Bible in their everyday life.
- 33 The Bible is the basis for what is taught.
- 34 By now you have learned many good things from the Bible.

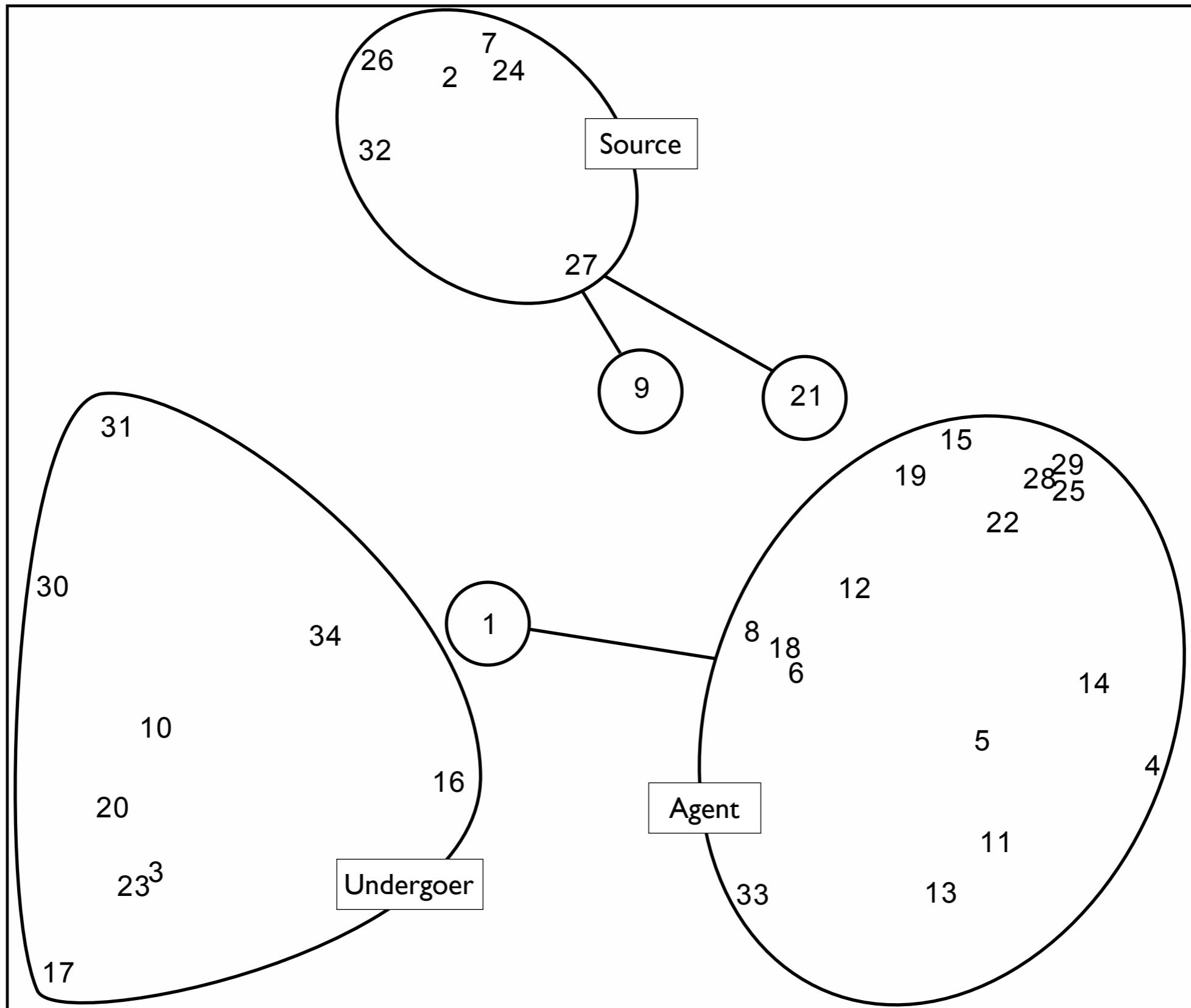
Albanian	Faroese	Estonian	Greenlandic
<i>bibla</i> Nominative	<i>biblian</i> Nominative	<i>þiibel</i> Nominative	<i>biibilip</i> Ergative
<i>biblën</i> Accusative	<i>bibliuna</i> Accusative	<i>þiiblit</i> Partitive	<i>biibli</i> Absolutive
<i>biblës</i> Genitive/Dative	<i>bibliunnar</i> Genitive	<i>þiibli</i> Genitive	<i>biibilmik</i> Instrumental
...	<i>bibliuni</i> Dative	<i>þiiblis</i> Inessive	<i>biibilmi</i> Locative
	...	<i>þiiblist</i> Elative	...
		...	

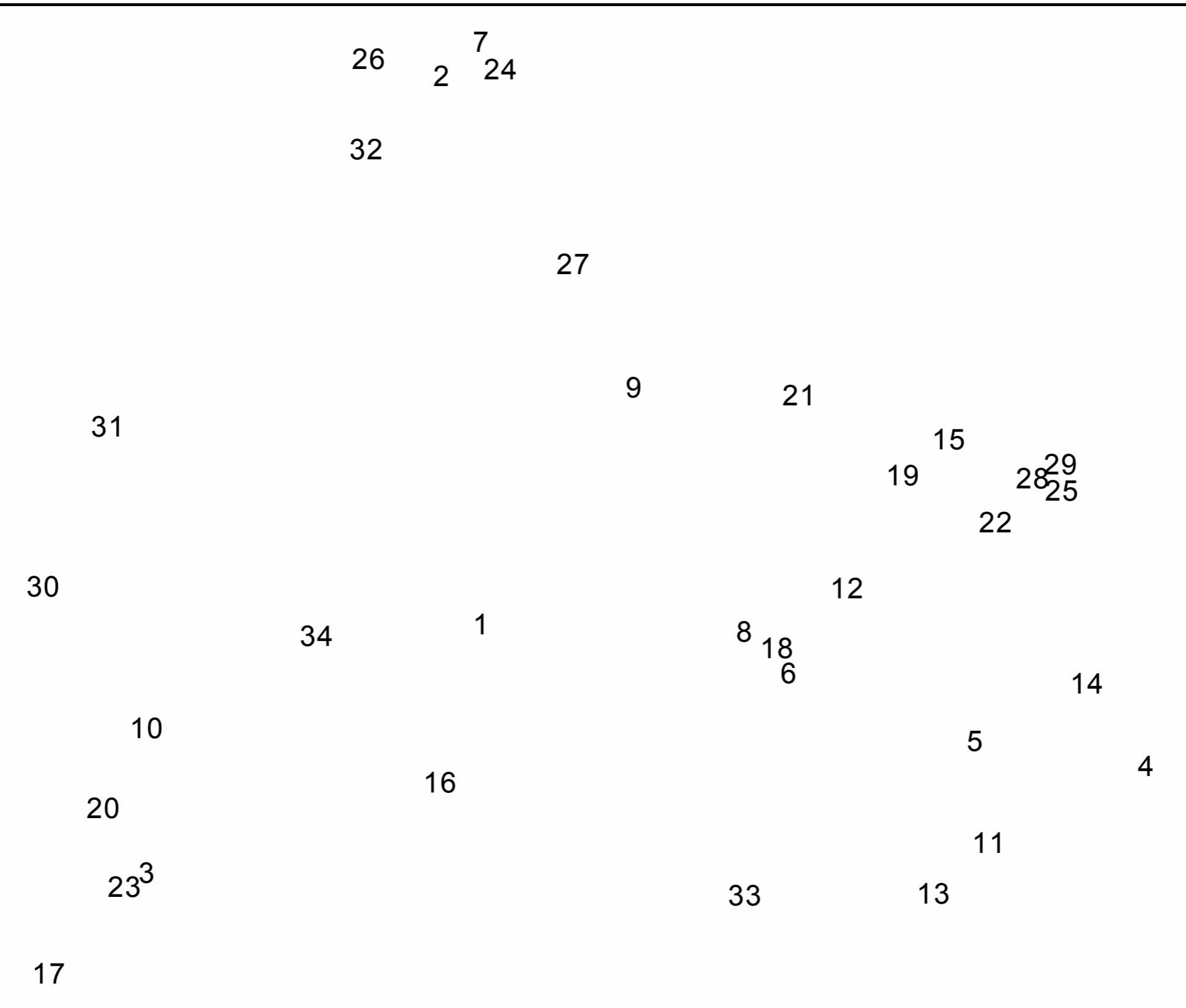
Context	Albanian	Faroese	Estonian	Greenlandic
1	bibla	bíbliuni	pübel	biibili
2	biblës	bíbliunnar	píibli	biibilimik
3	biblën	bíbliuna	píiblit	biibili
4	bibla	bíblian	pübel	biibili
5	bibla	bíblian	pübel	biiblip
6	bibla	bíbliuna	píibli	biibili
7	biblës	bíbliunnar	píibli	biibilimut
8	bibla	bíblian	píiblit	biiblip
9	bibla	NA	píiblit	biibili
10	biblën	bíbliuna	píiblit	biibilimik
11	bibla	bíblian	pübel	biibilimili
12	bibla	bíblian	pübel	biibilili
13	bibla	bíblian	pübel	biibilimi
14	bibla	bíblian	pübel	biibilimi
15	bibla	bíblian	pübel	biibilimi
16	bibla	bíbliuni	píibli	biibililu
17	biblën	bíbliuna	píiblit	biibilimik
18	bibla	bíblian	píiblis	biiblip
19	bibla	bíblian	píiblis	biibilimi
20	biblën	bíbliuna	píiblit	biibilimik
21	NA	bíblian	pübel	biibilimi
22	bibla	bíbliuni	pübel	biibili
23	biblën	bíbliuna	píiblit	biibilimillu
24	biblike	bíblian	píibli	biibilimi
25	bibla	bíblian	pübel	biibilimi
26	biblës	bíbliunnar	píibli	biibilimi
27	bibla	bíblian	píiblis	biibilimi
28	bibla	bíblian	pübel	biibilimi
29	bibla	bíblian	pübel	biibilimi
30	biblën	bíbliuna	píiblist	biibilimik
31	biblën	bíbliuni	píibli	biibilimik
32	biblës	bíbliuni	píibli	biibili
33	bibla	bíbliuna	pübel	biibilimik
34	bibla	bíbliuni	píiblist	biibilimeersunik

0,00	0,67	0,50	0,75	0,67	0,75	0,69	0,75	0,73	0,56	0,44	0,63	0,53	0,75	0,53	0,56	0,63	0,50	0,63	0,63	0,60	0,50	0,56	0,69	0,75	0,57	0,44	0,69	0,56	0,63	0,63	0,50	0,75	0,63
0,67	0,00	0,73	0,80	0,73	0,67	0,20	0,73	0,64	0,60	0,80	0,67	0,93	0,73	0,57	0,80	0,73	0,73	0,73	0,67	0,50	0,67	0,73	0,40	0,60	0,15	0,53	0,67	0,67	0,60	0,60	0,38	0,73	0,80
0,50	0,73	0,00	0,88	0,80	0,69	0,81	0,75	0,80	0,25	0,75	0,75	0,73	0,88	0,80	0,69	0,13	0,69	0,81	0,19	0,73	0,81	0,06	0,75	0,88	0,64	0,69	0,88	0,88	0,44	0,63	0,64	0,75	0,69
0,75	0,80	0,88	0,00	0,40	0,31	0,81	0,56	0,53	0,88	0,44	0,44	0,40	0,13	0,47	0,75	0,94	0,63	0,63	0,94	0,53	0,44	0,94	0,88	0,25	0,93	0,81	0,31	0,44	1,00	1,00	0,79	0,50	0,94
0,67	0,73	0,80	0,40	0,00	0,60	0,80	0,60	0,79	0,80	0,47	0,47	0,57	0,47	0,50	0,73	0,87	0,53	0,60	0,87	0,57	0,53	0,80	0,80	0,47	0,77	0,67	0,47	0,47	0,93	1,00	0,85	0,53	0,87
0,75	0,67	0,69	0,31	0,60	0,00	0,69	0,63	0,53	0,75	0,63	0,63	0,67	0,44	0,67	0,81	0,75	0,75	0,88	0,60	0,56	0,75	0,81	0,56	0,79	0,81	0,63	0,63	0,81	0,81	0,71	0,63	0,94	
0,69	0,20	0,81	0,81	0,80	0,69	0,00	0,63	0,60	0,75	0,81	0,75	0,93	0,75	0,67	0,81	0,81	0,81	0,81	0,60	0,75	0,81	0,50	0,75	0,29	0,63	0,69	0,69	0,69	0,50	0,81	0,81		
0,75	0,73	0,75	0,56	0,60	0,63	0,63	0,00	0,47	0,88	0,56	0,50	0,67	0,63	0,53	0,75	0,75	0,63	0,69	0,81	0,80	0,69	0,75	0,75	0,63	0,75	0,75	0,86	0,81	0,75				
0,73	0,64	0,80	0,53	0,79	0,53	0,60	0,47	0,00	0,73	0,80	0,73	0,86	0,67	0,64	0,80	0,87	0,87	0,87	0,80	0,86	0,67	0,87	0,73	0,67	0,69	0,73	0,67	0,73	0,80	0,80	0,62	0,80	0,80
0,56	0,60	0,25	0,88	0,80	0,75	0,75	0,88	0,73	0,00	0,81	0,75	0,80	0,81	0,80	0,69	0,25	0,69	0,81	0,19	0,67	0,88	0,25	0,69	0,81	0,57	0,63	0,81	0,81	0,44	0,63	0,57	0,63	0,81
0,44	0,80	0,75	0,44	0,47	0,63	0,81	0,56	0,80	0,81	0,00	0,56	0,27	0,38	0,27	0,69	0,81	0,44	0,44	0,81	0,53	0,31	0,75	0,81	0,38	0,86	0,63	0,31	0,19	0,88	0,88	0,86	0,56	0,81
0,63	0,67	0,75	0,44	0,47	0,63	0,75	0,50	0,73	0,75	0,56	0,00	0,53	0,50	0,47	0,69	0,81	0,44	0,56	0,81	0,33	0,63	0,75	0,63	0,50	0,64	0,63	0,56	0,56	0,88	0,88	0,71	0,69	0,81
0,53	0,93	0,73	0,40	0,57	0,67	0,93	0,67	0,86	0,80	0,27	0,53	0,00	0,27	0,36	0,60	0,73	0,40	0,33	0,80	0,43	0,40	0,73	0,73	0,33	0,77	0,60	0,33	0,33	0,80	0,80	0,85	0,60	0,73
0,75	0,73	0,88	0,13	0,47	0,44	0,75	0,63	0,67	0,81	0,38	0,50	0,27	0,00	0,33	0,69	0,88	0,56	0,50	0,88	0,40	0,44	0,88	0,75	0,13	0,79	0,69	0,19	0,31	0,94	0,94	0,79	0,44	0,88
0,53	0,57	0,80	0,47	0,50	0,67	0,67	0,53	0,64	0,80	0,27	0,47	0,36	0,33	0,00	0,73	0,87	0,33	0,27	0,80	0,50	0,20	0,80	0,53	0,20	0,57	0,33	0,20	0,07	0,80	0,80	0,64	0,60	0,73
0,56	0,80	0,69	0,75	0,73	0,81	0,81	0,75	0,80	0,69	0,69	0,69	0,60	0,69	0,73	0,00	0,63	0,63	0,75	0,69	0,80	0,75	0,69	0,75	0,75	0,71	0,75	0,75	0,81	0,69	0,56	0,57	0,63	0,44
0,63	0,73	0,13	0,94	0,87	0,75	0,81	0,75	0,87	0,25	0,81	0,81	0,73	0,88	0,87	0,63	0,00	0,75	0,88	0,19	0,80	0,94	0,13	0,81	0,94	0,71	0,75	0,94	0,94	0,31	0,50	0,79	0,63	0,63
0,50	0,73	0,69	0,63	0,53	0,75	0,81	0,63	0,87	0,69	0,44	0,44	0,40	0,56	0,33	0,63	0,75	0,00	0,25	0,75	0,53	0,44	0,69	0,63	0,56	0,64	0,38	0,56	0,44	0,81	0,81	0,64	0,69	0,69
0,63	0,73	0,81	0,63	0,60	0,75	0,81	0,69	0,87	0,81	0,44	0,56	0,33	0,50	0,27	0,75	0,88	0,25	0,00	0,81	0,47	0,31	0,81	0,56	0,38	0,57	0,31	0,38	0,25	0,81	0,81	0,64	0,69	0,75
0,63	0,67	0,19	0,94	0,87	0,88	0,81	0,81	0,80	0,19	0,81	0,81	0,80	0,88	0,80	0,69	0,19	0,75	0,81	0,00	0,80	0,88	0,19	0,75	0,81	0,64	0,69	0,69	0,81	0,63	0,64	0,69	0,69	
0,60	0,50	0,73	0,53	0,57	0,60	0,60	0,80	0,86	0,67	0,53	0,33	0,43	0,40	0,50	0,80	0,80	0,53	0,47	0,80	0,00	0,60	0,73	0,40	0,40	0,46	0,53	0,47	0,47	0,80	0,80	0,62	0,67	0,87
0,50	0,67	0,81	0,44	0,53	0,56	0,75	0,69	0,67	0,88	0,31	0,63	0,40	0,44	0,20	0,75	0,94	0,44	0,31	0,88	0,60	0,00	0,88	0,75	0,31	0,71	0,56	0,31	0,19	0,88	0,81	0,57	0,56	0,75
0,56	0,73	0,06	0,94	0,80	0,75	0,81	0,75	0,87	0,25	0,75	0,75	0,73	0,88	0,80	0,69	0,13	0,69	0,81	0,19	0,73	0,88	0,00	0,75	0,88	0,64	0,69	0,88	0,88	0,44	0,63	0,71	0,75	0,69
0,69	0,40	0,75	0,88	0,80	0,81	0,50	0,75	0,73	0,69	0,81	0,63	0,73	0,75	0,53	0,75	0,81	0,63	0,56	0,75	0,40	0,75	0,75	0,00	0,63	0,21	0,38	0,63	0,63	0,69	0,63	0,43	0,81	0,75
0,75	0,60	0,88	0,25	0,47	0,56	0,75	0,63	0,67	0,81	0,38	0,50	0,33	0,13	0,20	0,75	0,94	0,56	0,38	0,81	0,40	0,31	0,88	0,63	0,00	0,64	0,56	0,06	0,19	0,88	0,88	0,64	0,50	0,81
0,57	0,15	0,64	0,93	0,77	0,79	0,29	0,79	0,69	0,57	0,86	0,64	0,77	0,79	0,57	0,71	0,71	0,64	0,57	0,64	0,46	0,71	0,64	0,21	0,64	0,00	0,36	0,64	0,64	0,57	0,31	0,86	0,71	
0,44	0,53	0,69	0,81	0,67	0,81	0,63	0,75	0,																									

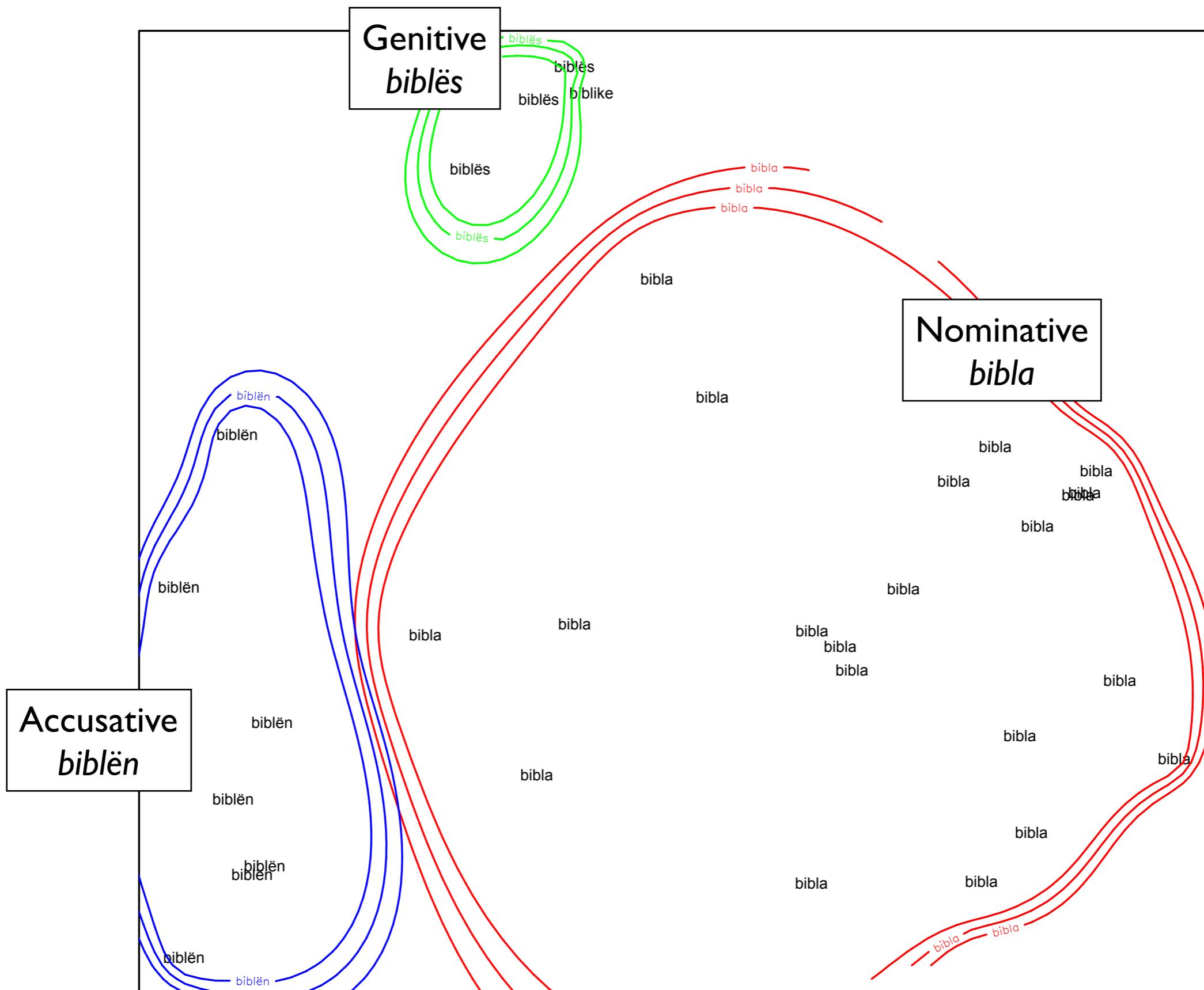




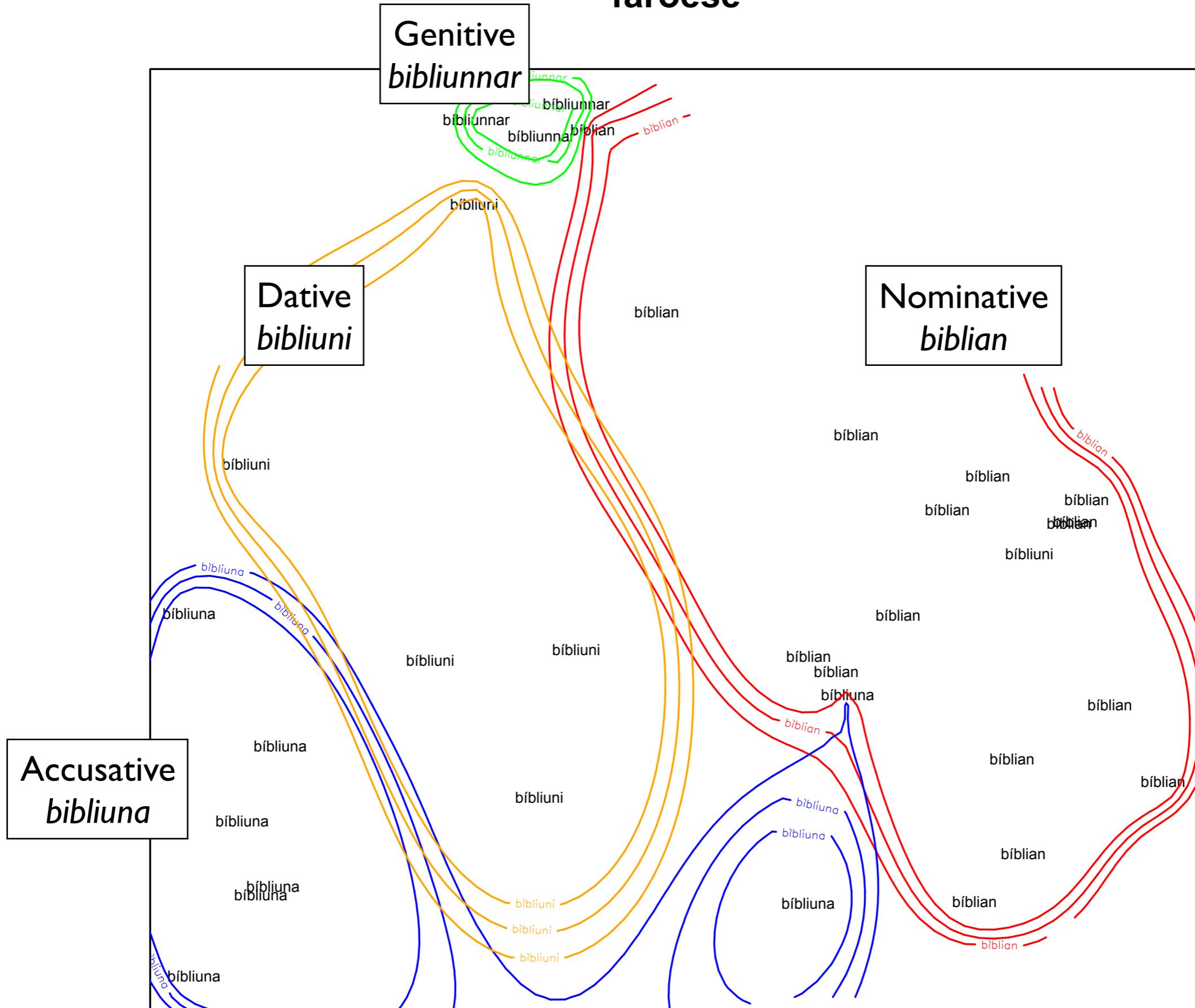




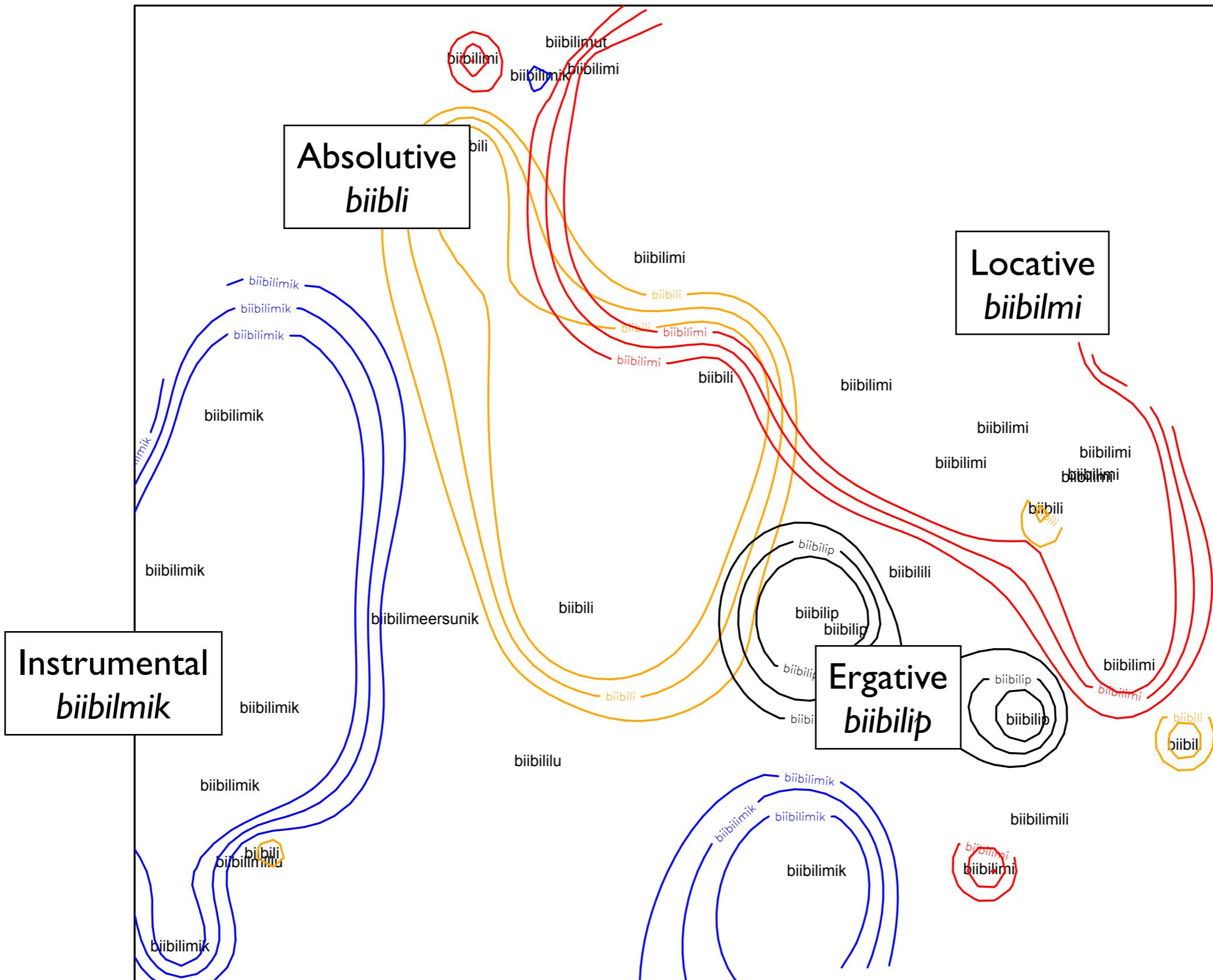
albanian



faroeese



greenlandic



Multi-alignment

- A. Biological Phylogenetics
(alignment of nucleotides/amino-acids)
- B. Historical Linguistics
(alignment of sounds in words)
- C. Functional Language Comparison
(alignment of words in sentences)

Three of a kind?

- Basically the same problem:
 - ▶ substrings within strings have to be aligned
- However, there are differences
 - ▶ the information of the substrings differs strongly
 - ▶ the lengths of the strings differs strongly

Parts/Length Ratio

(“informativity of the parts”)

- DNA: order 10^{-3} (viz. $\sim 4 / 1000s$)
(nucleotides vs. length of strings)
- Protein: order 10^{-1} (viz. $\sim 20 / 100s$)
(amino acids vs. length of protein)
- Words: order 10^{+1} (viz. $\sim 40 / 6$)
(sounds vs. length of word)
- Sentences: order 10^{+3} (viz. $\sim 1000s / 10$)
(number of words vs. length of sentence)

A different view of multi-alignment

- Multi-alignment is mostly seen as the problem of adding gaps in the right places
- Crossing alignments (“metathesis”) is seen as a nuisance / special case
- Now: words in sentences have massive crossing alignments.

A different view of multi alignment

- Consider multi-alignment as constrained partitioning (“flat clustering”)
- Partitioning
 - ▶ take all elements in all species/languages, and put them into groups (“mulit-alignments”)
- Constraints, e.g.
 - ▶ prefer groups across species/languages
 - ▶ ordering can be more or less strongly obeyed