Identifying Textual Clusters with Non-negative Matrix Factorization

Joey McCollum* 17 September 2020

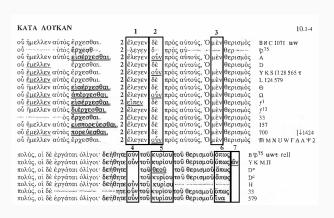
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· Start with collation—aligning their texts at variation units

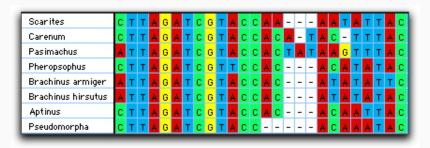
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KATA AOTKAN
                                                                                    10.1-4
οῦ ήμελλεν αύτὸς ἔρχεσθαι.
                               2 έλεγεν δὲ πρὸς αύτούς. Ὁ μὲν θερισμὸς
                                                                        B R C 1071 uw
ού ..... υτὸς ἔρχεσθ.....
                               2 -λεγεν δ. πρός αύ...... θεο......
                                                                         2375
ού ήμελλεν αύτὸς εισέρχεσθαι.
                               2 έλεγεν ούν πρός αὐτούς, Ό μὲν θερισμός
ດນີ້ ຂຶ້ນຂໍλλεν
                ἔργεσθαι.
                               2 ἔλεγεν δὲ πρὸς αὐτούς, Ὁ
                                                              θερισμός
οὖ ἔμελλεν αύτὸς ἔρχεσθαι.
                               2 έλεγεν ούν ποὸς αὐτούς. Ό μὲν θεοισμὸς
                                                                         Y K S IT 28 565 T
ού εμελλεν αυτός εργεσθαι.
                               2 έλεγεν δὲ πρὸς αὐτούς, Ὁ μὲν θερισμὸς
                                                                        L 124 579
ού ήμελλεν αύτὸς εἰσέργεσθαι.
                               2 έλεγεν δὲ πρὸς αὐτούς, Ὁ μὲν θερισμὸς
ού ἔμελλεν αύτὸς ἀπέργεσθαι.
                               2 έλεγεν ούν πρὸς αὐτούς, Ὁ μὲν θερισμὸς
                                                                         Ω
ού εμελλεν αύτὸς εἰσέργεσθαι.
                               2 είπεν δὲ πρὸς αὐτούς, Ὁ μὲν θερισμὸς
                                                                        f1
ού ήμελλεν αύτὸς διέρχεσθαι.
                               2 έλεγεν δὲ πρὸς αὐτούς. Ὁ μὲν θερισμὸς
ος ἔρχεσθαι.
                               2 έλεγεν δὲ πρὸς αὐτούς, Ὁ μὲν θερισμὸς
ού ημελλεν αύτὸς εισπορεύεσθαι, 2 έλεγεν δὲ πρὸς αύτούς, Ο μὲν θερισμὸς
                                                                        157
ού εμελλεν αύτός πορεύεσθαι.
                               2 έλεγεν δὲ πρὸς αὐτούς. Ὁ μὲν θερισμὸς
                                                                        700
                                                                                    Г↓1424
                               2 έλεγεν ούν πρός αύτούς, Ο μέν θερισμός
ού ήμελλεν αύτὸς ἔργεσθαι.
                                                                        TRMNUWFAA 42
πολύς, οι δὲ ἐργάται όλίγοι δεήθητε ούν τοῦ κυρίου τοῦ θερισμοῦ ὅπως
                                                                     B 9\75 HW7 rell
πολύς, οι δὲ ἐργάται όλίγοι δεήθητε ούν τοῦ κυρίου τοῦ θερισμοῦ ὅπως ἀν
                                                                     YKMII
πολύς, οἱ δέ έργάται όλίγοι δεήθητε
                                     τοῦ θεοῦ τοῦ θερισμοῦ ὅπως
                                                                     n*
πολύς, οί δε εργάται όλίνοι δεήθητε τοῦ κυρίου τοῦ θερισμοῦ όπως
                                                                     DC
πολύς, οί δε εργάται όλίνοι: δεήθητε ούν τού κυρίου .....
                                                                      Н
πολύς, οι δὲ ----- ητε ούν τοῦ κυρίου τοῦ θερισμοῦ ὅπως
                                                                     33
πολύς, οι δὲ ἐργάται όλίνοι: δεήθητε ούν τοῦ κυρίου τοῦ θερισμοῦ ἴνα
                                                                     579
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(Source: Swanson, New Testament Greek Manuscripts, Luke, 183)

· Start with collation—aligning their texts at variation units



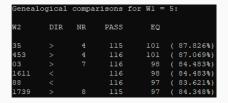
- · Comparable to DNA sequence alignment¹
 - manuscripts ←→ taxa / species
 - variation units ←→ sites
 - variant readings ←→ bases (A, C, G, T) and gaps (–)



(Source: http://www.sequence-alignment.com)

^{1.} For the fascinating history of this relationship, see Lin, *The Erotic Life of Manuscripts*.

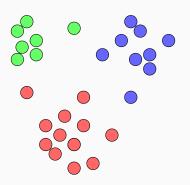
- This provides a simple basis of comparison between pairs of manuscripts
 - · Number of units where both agree
 - For a proportion, divide by number of units where the readings of both are known
- "Pre-genealogical coherence" in the Coherence-Based Genealogical Method (CBGM)



 Can we use mutual agreement to classify manuscripts into groups?

The Quantitative Method

- Colwell and Tune: if manuscripts agree significantly more with one another than they do with other manuscripts, then they form a family, or text-type²
 - $\cdot \geq$ 70% with one another, and 10% more than with others

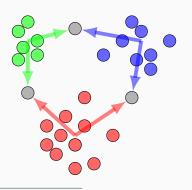


2. Colwell and Tune, "Quantitative Relationships."

The Quantitative Method

· Problems:

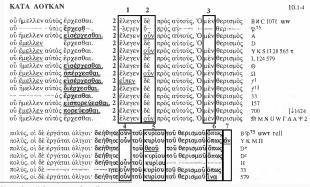
- All units (including those involving singular readings and common scribal errors) have equal weight
- · Mixture in the transmission process is a problem³



3. Epp, "Textual Clusters."

The Quantitative Method

- For efficiency and accuracy, comparisons should be done on the basis of informative points of variation⁴
- · Specific readings, not the variation units containing them
- · But how do we know which ones are the most informative?



4. Colwell, "Method in Locating."

The Claremont Profile Method

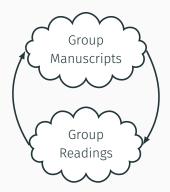
- Start with an established set of manuscript groups⁵
- Filter out variation units involving common types of variation and singular / subsingular readings to get a set of test passages
- Readings supported by group manuscripts = the group's profile



^{5.} Wisse, Profile Method.

The Claremont Profile Method

- This allows us to isolate informative readings for group classification
- · Also robust to mixture
- But it needs manuscript groups to be established first!
- "Good manuscripts have good readings, and good readings are found in good manuscripts"



- Non-negative matrix factorization (NMF), a machine learning technique, uses this circular relationship to solve both problems
- Represent our collation as a matrix A with a row for each variant reading and a column for each manuscript
- m rows by n columns

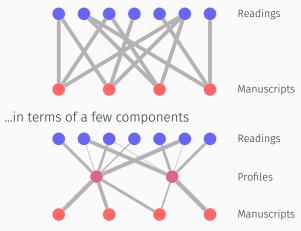
				_	_			
		\mathfrak{P}^{75}	Α	В	D	K	f^1	579
Unit 1	ἔλεγεν	1	1	1	1	1	0	1
Omt 1	εἶπεν	0	0	0	0	0	1	0
Unit 2	δè	1	0	1	1	0	1	1
OIIIt 2	οὖν	0	1	0	0	1	0	0
Unit 3	μὲν	0	1	1	0	1	1	1
Omt 3	omit	0	0	0	1	0	0	0
Unit 4	οὖν	ν 1 1 1 0 1	1	1				
OIIIt 4	omit	0	0	0	1 0 0 0	0		
Unit 5	κυρίου	ρίου 1 1 1 0 1 1	1					
Omt 3	θεοῦ	0	0	0	1	0	0	0
Unit 6	őπως	1	1	1	1	1	1	0
Omto	ἵνα	0	0	0	0	0	0	1
Unit 7	omit 1 1 1 1 0 1	1	1					
Onit /	ầν	0	0	0	0	1	0	0

 The goal is to approximate this original matrix as the product of two smaller matrices:

$A \approx WH$

- Specify a number k of underlying textual profiles (there are metrics for finding good choices)
- W: m rows and k columns; defines group readings
- **H**: *k* rows and *n* columns; defines makeup of manuscripts in terms of profiles

• "High-fidelity data compression"; reconstruct the original data...



· A "sum of parts" model that is robust to mixture

- The process:
 - 1. Start with guesses for \mathbf{W} and \mathbf{H}
 - 2. Fix \mathbf{W} , optimize the weights in \mathbf{H} (Quantitative Method)
 - 3. Fix ${f H}$, optimize the weights in ${f W}$ (Claremont Profile Method)
 - 4. Repeat steps 2 and 3 until the difference between ${\bf A}$ and ${\bf WH}$ no longer decreases



• Guaranteed to terminate with *locally optimal* groupings in ${f W}$ and ${f H}^6$

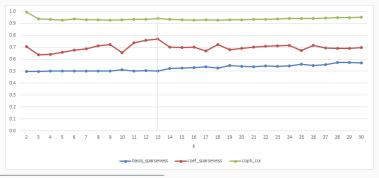
^{6.} Grippo and Sciandrone, "On the Convergence."

- Tommy Wasserman's collation of Jude contains 1346 variant readings and 560 manuscripts (including lectionaries)⁷
- Filtering out 42 fragmentary manuscripts (< 300 known readings) yields a matrix $\bf A$ with m=1346 rows and n=518 columns
- The fragmentary manuscripts can be classified after groups are established⁸

^{7.} Wasserman, The Epistle of Jude.

^{8.} For details, see the appendix of McCollum, "Biclustering Readings and Manuscripts."

- We select the number of profiles *k* based on several factors:
 - · Overlap of readings in profiles
 - · Mixture of profiles in manuscripts
 - Consistency of manuscript groupings when random starting points are used (the *cophenetic correlation coefficient*)⁹



9. Brunet et al., "Metagenes and Molecular Pattern Discovery."

• The k = 13 groups identified by NMF correspond to groups in the Catholic Epistles identified in the literature

Members (by Gregory-Aland number)	Group
920, 1277, 1859, 1719, 452, 1857, 1871, 941,	K (von Soden)
1103, 1352, etc.	
141, 204, 394, 444, 1101, 1723, 1737, 1752,	K ^r (von Soden)
1865, 2221, etc.	
390, 1863, 912, 234, 1861, 2085, 1753, 2279, 42,	K ^c (von Soden)
996, etc.	
L606, L938, L145, L840, L740, L2106, L2394,	Lectionary (Colwell)
L809, L1279, L62, etc.	
606, 454, 641, 103, 221, 2125, 314, 250, 1888,	O, Θδ Commentaries
393, etc.	(von Soden)
619, 1780, 1175, 330, 1769, 2516, 917, 451,	f^{1780} (unidentified)
1162, 601, etc.	
1563, 1718, 1425, 1359, 1066, 0142, 056	f ⁰¹⁴² (unidentified)

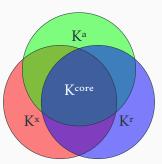
• The k = 13 groups identified by NMF correspond to groups in the Catholic Epistles identified in the literature

Members (by Gregory-Aland number)	Group	
03, 623, \mathfrak{P}^{72} , 81, 5, 326, 33, 1837, 93, 665, etc.	H (von Soden)	
321, 918, 307, 453, 2197, 2818, 1678, 94, 2186,	f453 (Spencer, Wachtel,	
1840, etc.	Howe)	
323, 1241, 322, 1739, 1881, 2298, 6	f ¹⁷³⁹ (Zuntz, Geer)	
1505, 2495, 1611, 1292, 630, 2200, 1765, 1832,	f ²¹³⁸ /Harklean	
2494, 876, etc.	(Amphoux)	
1843, 1869, 506, 1903, 489, 927, 203, 1868,	l (von Soden)	
1729, 1873, etc.		
915, 88, 459, 104, 1846, 1838, 1842, 1845	f ⁹¹⁵ (unidentified)	

- Applying NMF to Morrill's collation of all continuous-text manuscripts of John 18 illustrates some of the idiosyncrasies of the method and how to deal with them¹⁰
- Significantly larger and more "square" collation: m=1545 variant readings and n=1610 manuscripts after filtering out fragmentary manuscripts (< 350 known readings)
- \cdot (Recall that the collation matrix for Jude was 1346 imes 518)

^{10.} Morrill, "Complete Collation and Analysis."

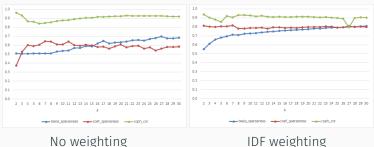
- Applying NMF to the matrix as-is separates readings common to multiple groups into their own "core" profiles
- No manuscripts belong to these profiles, but many appear "mixed" with it
- "Primary readings" vs. "Secondary readings" in profiles
- Symptom of volume and similarity of manuscripts, especially Byzantine ones



• To remedy this, weigh readings in the original matrix by their inverse document frequency (IDF)11

$$\log \frac{n}{\#\{\text{MSS with reading}\}}$$

- Removing singular readings is helpful in this setting
- Encourages NMF to isolate unique group readings in profiles



^{11.} Jones, "Statistical Interpretation."

IDF weighting

 \cdot With k=12, NMF identifies known groups from the literature

Members (by Gregory-Aland number)	Group
2605, 492, 1215, 2897, 1090, 1567, 1210, 851,	K ^x (von Soden)
494, 2406, etc.	
47, 1126, 61, 1138, 58, 56, 189, 1236, 825, 1614,	K ^r (von Soden)
etc.	
2902, 1219, 1079, 489, 114, 2404, 389, 2193,	Ka (von Soden)
699, 1627, etc.	
1534, 741, 857, 744, 2735, 1160, 817, 1261,	Θε Commen-
2470, 833, etc.	taries (von
	Soden)
892, 977, 555, 16, 152, 513, 1243, 829, 348, 1579,	$f^{16}+f^{1216}$
etc.	(Wisse)
1663, 1413, 2291, 86, 569, 71, 1170, 1014, 1531,	M27+Cl1531
2705, etc.	(Wisse)

• With k = 12, NMF identifies known groups from the literature

Members (by Gregory-Aland number)	Group
01, 032, 05, 579, 1654, 2561, 1242	Egyptian
1820, 2129, 865, 033, 019, 1819, 213, 03, 33,	Alexandrian
1321, etc.	
1, 1582, 357, 138, 565, 209, 994, 2713, 2575,	f1 (Lake)
1784, etc.	
13, 788, 826, 828, 543, 69, 346, 1689, 124, 2786,	f13 (Lake and Lake,
etc.	Geerlings)
2524, 1001, 1268, 2397, 352, 2728, 132, 175,	Cl1001+Cl352 (Wisse)
1701, 2252, etc.	
1446, 1050, 706, 1457, 827, 2620, 1128, 0211,	Cl827 (Wisse)
2707, 1402, etc.	

Concluding Observations

- In John 18, Gregory-Aland 03 (Codex Vaticanus, B) stands out as an instructive example
- Appears to be mixed between the "Egyptian" and "Alexandrian" profiles, but could preserve a text earlier than both

	"03"
Kx	0.0290
Cl1001+Cl352	0.0000
Theophylact	0.0000
f13	0.0000
f1	0.0000
Alexandrian	1.1248
Egyptian	1.0556
Kr	0.0000
Ka	0.0000
M27+Cl1531	0.0000
f16+f1216	0.0000
CI827	0.0000

- · NMF identifies relationships, but not their directions
- Pre-genealogical, but not genealogical



Concluding Observations

- · The advantage: few assumptions are required
- Intended for use in "pre-processing" (manuscript and test reading selection)
- · Useful for other applications (new manuscript classification)
- Work in progress: applying NMF to ~2000 manuscripts in the pericope adulterae (with Maurice A. Robinson)

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