

SYNTAX Score Management

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Introduction: Syntax Score:

Synthesis of Multiple Coronary Lesion Complexity Scoring Systems:

- **Coronary tree segments based on AHA classification modified for the ARTS I&II studies** (*Circulation* 1975; 51:31-3 & *Semin Interv Cardiol* 1999; 4:209-19)
- **Modified Leaman score** (*Circulation* 1981; 63: 285-292)
- **ACC/AHA lesions classification system** (*Circulation* 2001;103:3019-3041)
- **MEDINA classification** (*Rev Esp Cardiol.* 2006;59(2):183-4)
- **Total occlusion classification** (*J Am Coll Cardiol*, 1997;30:649-56)

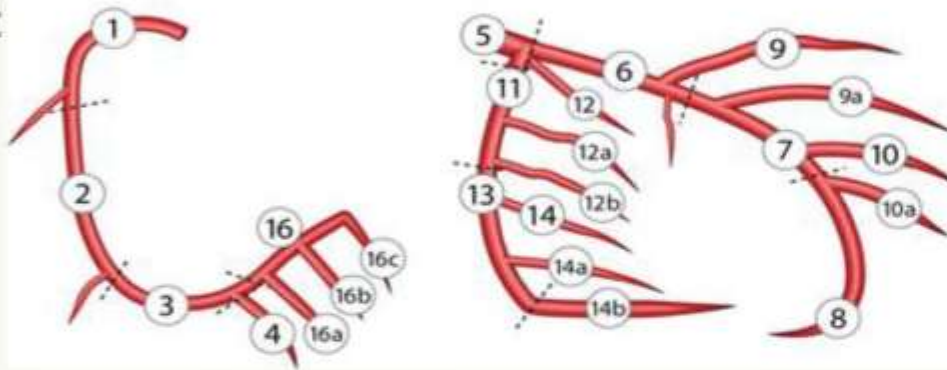
Syntax Score calculator (11 Questions)

- **Q1: Dominance**
- **Q2: Specify diseased segment numbers**
- **Q3: Total Occlusion**
- **Q4: Trifurcation**
- **Q5: Bifurcation**
- **Q6: Aorta Ostial**
- **Q7: Severe Tortuosity**
- **Q8: Length**
- **Q9: Heavy Calcification**
- **Q10: Thrombus**
- **Q11: Diffusely diseased and narrowed segments**

Syntax Score tutorial

Q1. Dominance

Right



Left

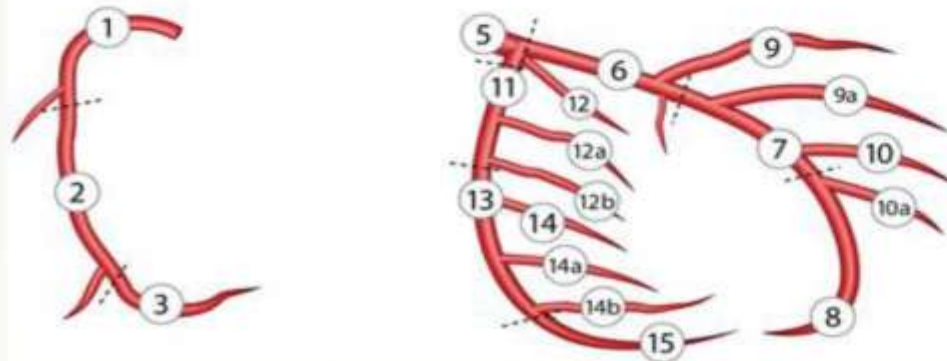


Table 1. Segment weighing factors

Segment No		Right dominance	Left dominance
1	RCA proximal	1	0
2	RCA mid	1	0
3	RCA distal	1	0
4	Posterior descending artery	1	n.a.
16	Posterolateral branch from RCA	0.5	n.a.
16a	Posterolateral branch from RCA	0.5	n.a.
16b	Posterolateral branch from RCA	0.5	n.a.
16c	Posterolateral branch from RCA	0.5	n.a.
5	Left Main	5	6
6	LAD proximal	3.5	3.5
7	LAD mid	2.5	2.5
8	LAD apical	1	1
9	First diagonal	1	1
9a	First diagonal ^a	1	1
10	Second diagonal	0.5	0.5
10a	Second diagonal ^a	0.5	0.5
11	Proximal circumflex artery	1.5	2.5
12	Intermediate/ anterolateral artery	1	1
12a	Obtuse marginal ^a	1	1
12b	Obtuse marginal ^b	1	1
13	Distal circumflex artery	0.5	1.5
14	Left posterolateral	0.5	1
14a	Left posterolateral ^a	0.5	1
14b	Left posterolateral ^b	0.5	1
15	Posterior descending	n.a.	1

**for definitions segment numbers see Tutorial (www.syntaxscore.com)*

Syntax Score: segment weight X multiplication factor

- Multiplication factor based on severity of stenosis
(specific % stenosis is *not* used)

Total Occlusion

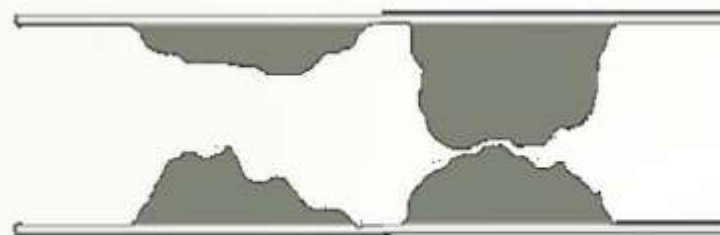
x 5



DS 100%

Significant lesion 50-99%

x 2



DS 50% - 99%

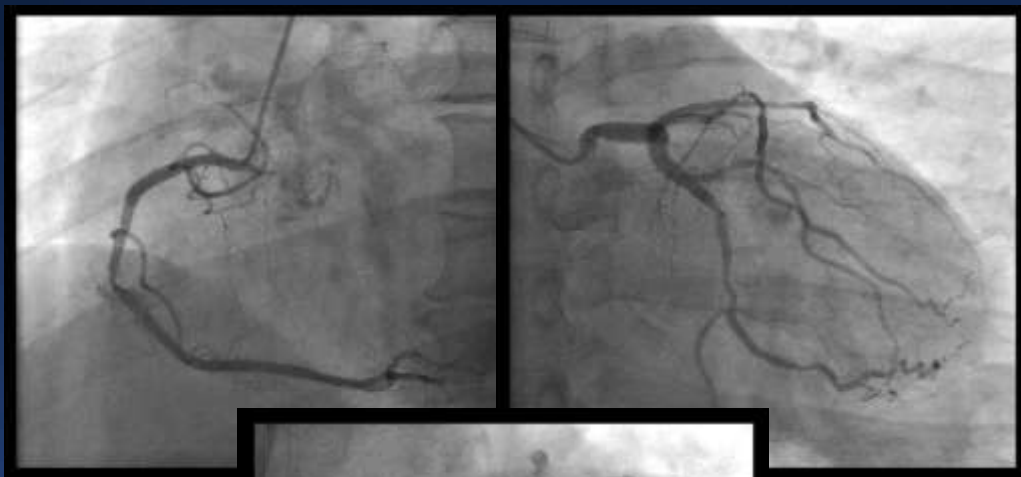


SYNTAX Score

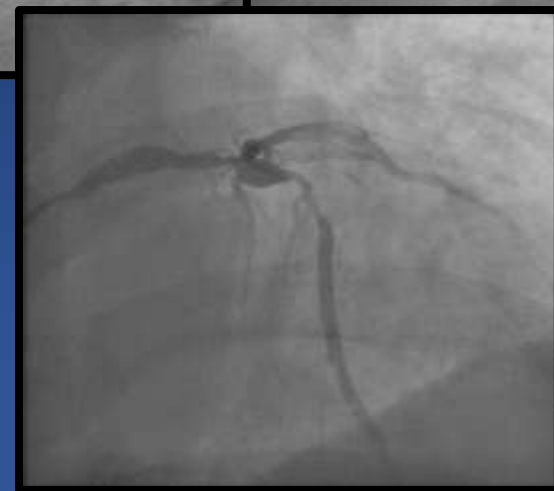


Why Should We Syntax Score?

Complexity Scoring System



2VD
SYNTAX Score : 10



3VD
SYNTAX Score : 54.5

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ESTABLISHED IN 1812

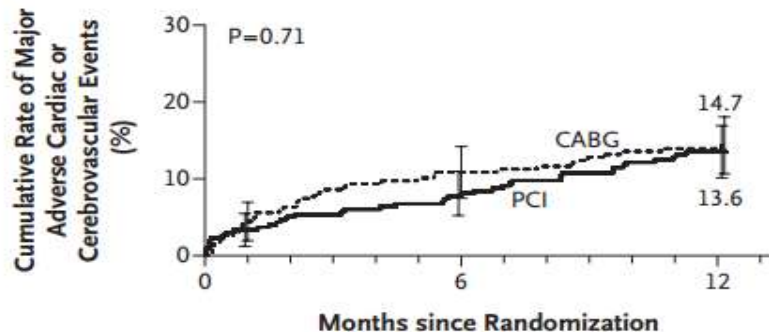
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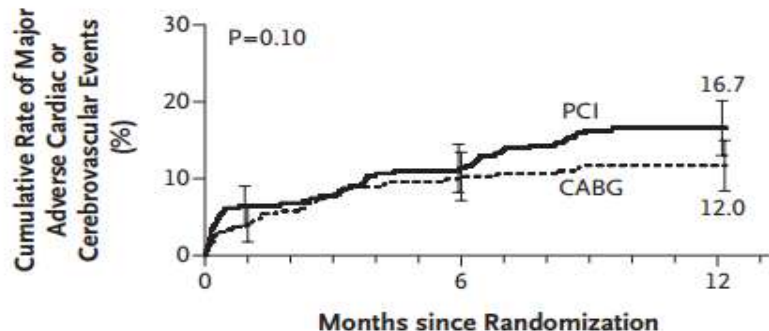
Percutaneous Coronary Intervention versus Coronary-Artery Bypass Grafting for Severe Coronary Artery Disease

Patrick W. Serruys, M.D., Ph.D., Marie-Claude Morice, M.D., A. Pieter Kappetein, M.D., Ph.D.,
Antonio Colombo, M.D., David R. Holmes, M.D., Michael J. Mack, M.D., Elisabeth Stähle, M.D.,
Ted E. Feldman, M.D., Marcel van den Brand, M.D., Eric J. Bass, B.A., Nic Van Dyck, R.N., Katrin Leadley, M.D.,
Keith D. Dawkins, M.D., and Friedrich W. Mohr, M.D., Ph.D., for the SYNTAX Investigators*

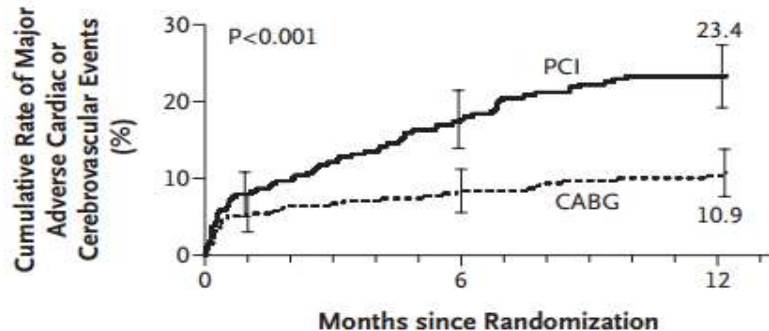
A Low SYNTAX Score



B Intermediate SYNTAX Score



C High SYNTAX Score



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Intervention versus Coronary-Artery Bypass Grafting for Severe Coronary Artery Disease

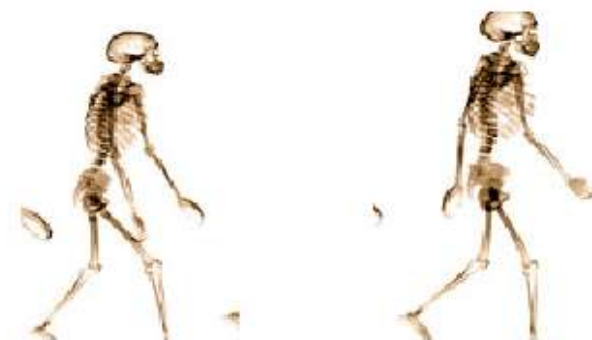
Claude Morice, M.D., A. Pieter Kappetein, M.D., Ph.D., Jacques Lesman, M.D., Michael J. Mack, M.D., Elisabeth Stähle, M.D., J. J. P. M. Bax, M.D., Eric J. Bass, B.A., Nic Van Dyck, R.N., Katrin Leadley, M.D., W. Mohr, M.D., Ph.D., for the SYNTAX Investigators*

Figure 3. Rates of Major Adverse Cardiac or Cerebrovascular Events among the Study Patients, According to Treatment Group and SYNTAX Score Category.

Kaplan-Meier curves are shown for the percutaneous coronary intervention (PCI) group and the coronary-artery bypass grafting (CABG) group for major adverse cardiac or cerebrovascular events at 12 months. The 12-month event rates were similar between the two treatment groups for patients with low SYNTAX scores (0 to 22) (Panel A) or intermediate SYNTAX scores (23 to 32) (Panel B). Among patients with high SYNTAX scores (≥33, indicating the most complex disease), those in the PCI group had a significantly higher event rate at 12 months than those in the CABG group. SYNTAX scores were calculated at the core laboratory. The I bars indicate 1.5 SE. P values were calculated with the use of the chi-square test.



Evolution of the SYNTAX Score...



EuroIntervention
SYNTAX Score

**Clinical
SYNTAX Score**
Circulation
Cardiovascular Interventions

AHJ
**Global Risk
Classification**

**JACC
Interventions**
Global Risk

**JACC
Imaging**
**Functional
SYNTAX Score**

ACC.12
**Logistic Clinical
SYNTAX Score**

**CABG SYNTAX
Score**
PCR PARIS 2012

**JACC
Interventions**
**SYNTAX
Myocardial
Infarction Score**

**JACC
Imaging**
**Compositional
SYNTAX Score**

EuroIntervention
**Residual
SYNTAX Score**

**Non-Invasive,
Logistic,
Functional,
CABG, Residual,
SYNTAX Score...**



PRECOMBAT SYNTAX Score - AMC



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Supplemental Table 2. Baseline Angiographic Characteristics of the Patients, According to Study Group. *

Random

Angiographic Characteristics	PCI (N=300)	CABG (N=300)	P value
Diseased vessels, N (%)			0.68
Left main only	27 (9.0)	34 (11.3)	
Left main plus single vessel disease	50 (16.7)	53 (17.7)	
Left main plus double vessel disease	101 (33.7)	90 (30.0)	
Left main plus triple vessel disease	122 (40.7)	123 (41.0)	
Bifurcation left main involvement	200 (66.9)	183 (62.2)	0.24
Heavy calcification of left main	17 (5.7)	14 (4.7)	0.58
Diameter stenosis of left main, N (%)			0.12
> 50% and ≤ 70%	160 (53.3)	141 (47.0)	
> 70%	140 (46.7)	159 (53.0)	
Right coronary artery disease, N (%)	149 (49.7)	159 (53.0)	0.41
Restenotic lesion, N (%)	1 (0.3)	2 (0.7)	0.56
Chronic total occlusion, N (%)	2 (0.7)	2 (0.7)	1.0
SYNTAX score, mean±SD §	24.4±9.4	25.8±10.5	0.09
SYNTAX score by tertiles, N (%)			0.75
Score ≤ 19	95 (32.9)	85 (31.6)	
Score > 19 and ≤ 29	105 (36.3)	93 (34.6)	
Score > 29	89 (30.8)	91 (33.8)	

Supplemental Table 8. Baseline Characteristics of the Patients, According to Treatment Group in the Registry Cohort.

Registry

Characteristics	PCI Registry (N=475)	CABG Registry (N=335)	P value
Angiographic characteristics			
Extent of disease vessel			<0.001
Left main only	74 (15.8)	2 (0.6)	
Left main plus single vessel disease	126 (27.0)	13 (4.1)	
Left main plus double vessel disease	171 (36.6)	39 (12.2)	
Left main plus triple vessel disease	96 (20.6)	265 (83.1)	
Involved location			0.22
Ostial and shaft involvement	158 (33.8)	96 (29.7)	
Distal bifurcation involvement	309 (66.2)	227 (70.3)	
Right coronary artery disease	156 (32.8)	287 (85.7)	<0.001
Restenotic lesion	8 (1.7)	1 (0.3)	0.41
Chronic total occlusion	16 (3.4)	5 (1.5)	0.10
EuroSCORE value	3.1±2.2	3.7±2.1	<0.001
SYNTAX score	22.5±9.9	37.8±11.9	<0.001
Postprocedural hospital stay, days	2.3±17.4	10.9±26.7	<0.001

Syntax Score - AMC



Random Trial



Balance Group



SYNTAX SCORE Calculator Live Demonstration

SYNTAX SCORE



Welcome to the SYNTAX Score website. The SYNTAX Score is a unique tool to score complexity of coronary artery disease. However, it is very important to use this new scoring tool correctly, hence, it is strongly recommended to complete the tutorial first.

TUTORIAL

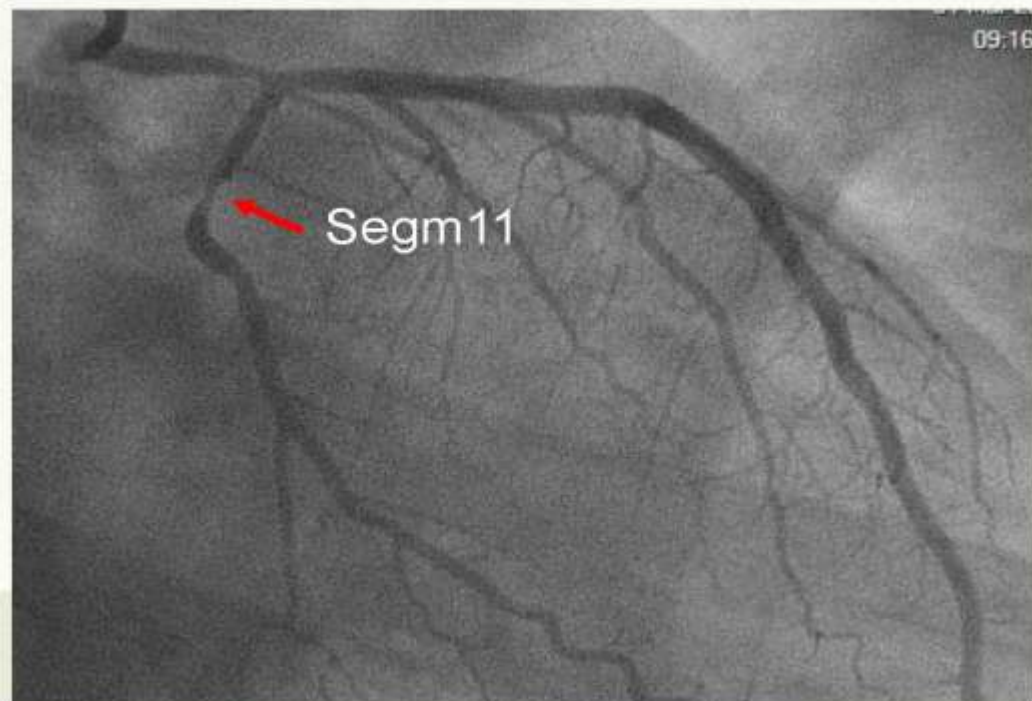
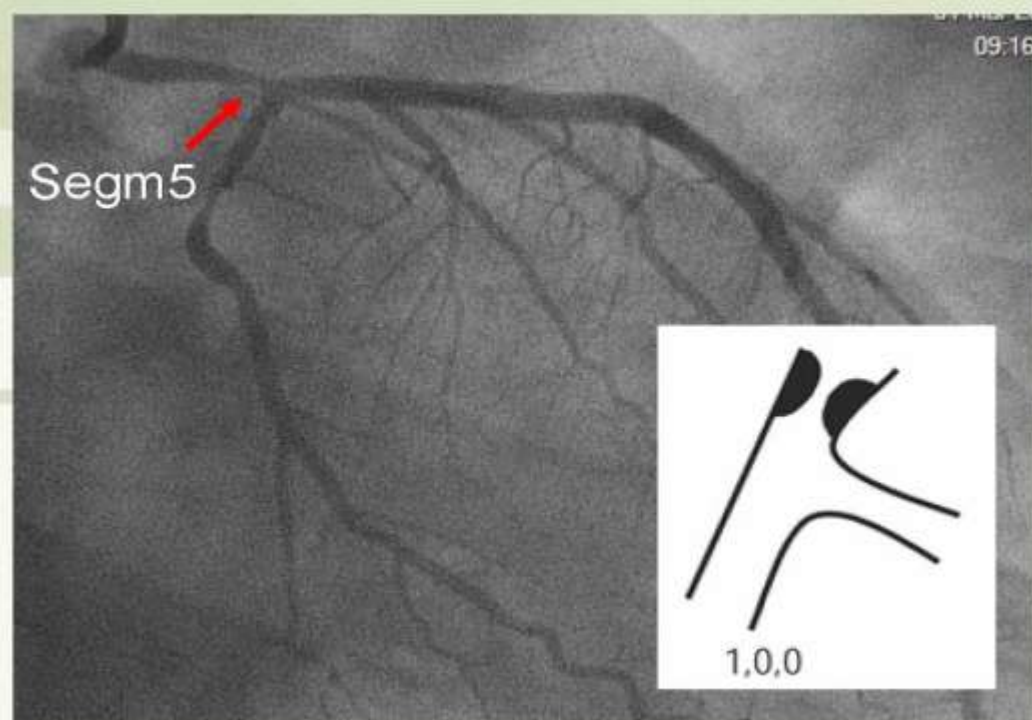
Knowledge of definitions is vital. Please use the tutorial prior to first calculator use.

[Start tutorial...](#)

CALCULATOR

Start using the calculator when you have successfully completed the tutorial.

[Start calculator...](#)



Total Syntax Score: 18

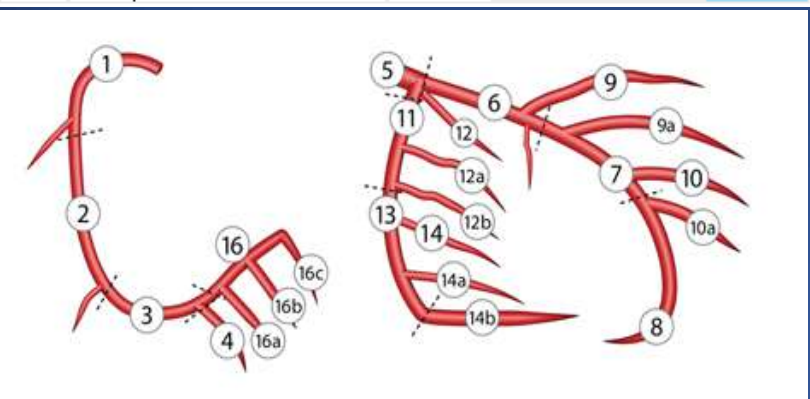
Lesion1: Left-main bifurcation (Medina 100)

Lesion2: Segment 11 (+ tortuosity)

Lesion3: Segment 1

SYNTAX SCORE Calculator

	Lesions:	1	2	3
Segments:				
RCA	RCA proximal	1	v	
	RCA mid	2		
	RCA distal	3		
	Posterior descending	4		
	Posterolateral from RCA	16		
	Posterolateral from RCA	16a		
LM	Left main	5		
	LAD proximal	6		
	LAD mid	7		
	LAD apical	8	v	
	First diagonal	9		
	Add. first diagonal	9a		
LAD	Second diagonal	10		
	Add. second diagonal	10a		
	Proximal circumflex	11		v
	Intermediate/anterolateral	12		
	Obtuse marginal	12a		
	Obtuse marginal	12b		
LCX	Distal circumflex	13		
	Left posterolateral	14		
	Left posterolateral	14a		
	Left posterolateral	14b		



4. Total occlusion (T.O.)

- a. ☐ No
b. ☐ Yes:

5. Trifurcation

- a. ☐ No
b. ☐ Yes

6. Bifurcation

- a. ☐ No
b. ☐ Yes

7. Aorto Ostial lesion

- a. ☐ No
b. ☐ Yes

8. Severe Tortuosity

- a. ☐ No
b. ☐ Yes

9. Length > 20 mm

- a. ☐ No
b. ☐ Yes

10. Heavy calcification

- a. ☐ No
b. ☐ Yes

11. Thrombus

- a. ☐ No
b. ☐ Yes

4. Total occlusion (T.O.)

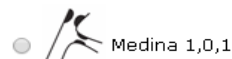
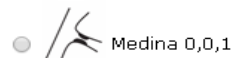
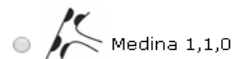
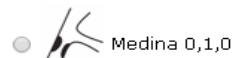
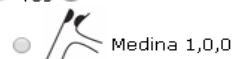
- a. ☐ No
b. ☐ Yes:

5. Trifurcation

- a. ☐ No
b. ☐ Yes

6. Bifurcation

- a. ☐ No
b. ☐ Yes



Bifurcation angulation (betw main vessel and side branch)

- a. ☐ No
b. ☐ Yes

8. Severe Tortuosity

- a. ☐ No
b. ☐ Yes

9. Length > 20 mm

- a. ☐ No
b. ☐ Yes

10. Heavy calcification

- a. ☐ No
b. ☐ Yes

11. Thrombus

- a. ☐ No
b. ☐ Yes

4. Total occlusion (T.O.)

- a. ☐ No
b. ☐ Yes:

I. Indicate the first segment number of the T.O.

LCX	Proximal circumflex	11	
-----	---------------------	----	--

II. is Age of T.O. > 3 months?

1. ☐ No
2. ☐ Yes
3. ☐ Unknown

III. Blunt stump?

1. ☐ No
2. ☐ Yes

IV. Bridging?

1. ☐ No
2. ☐ Yes

V. Specify the first segment number beyond the total occlusion that visualized by **antegrade or retrograde** contrast.

	Segment numbers:	Segment visualized by contrast:
	none	<input type="radio"/>
If, T.O. in segment 11	11	<input type="radio"/>
	13	<input type="radio"/>
	14	<input type="radio"/>

VI. Is there a sidebranch at the origin of the occlusion?

1. ☐ No
2. ☐ Yes, all sidebranches <1.5mm
3. ☐ Yes, all sidebranches >=1.5mm
4. ☐ Yes, both sidebranches <1.5mm and >=1.5mm are involved

7. Aorto Ostial lesion

- a. ☐ No
b. ☐ Yes

8. Severe Tortuosity

- a. ☐ No
b. ☐ Yes

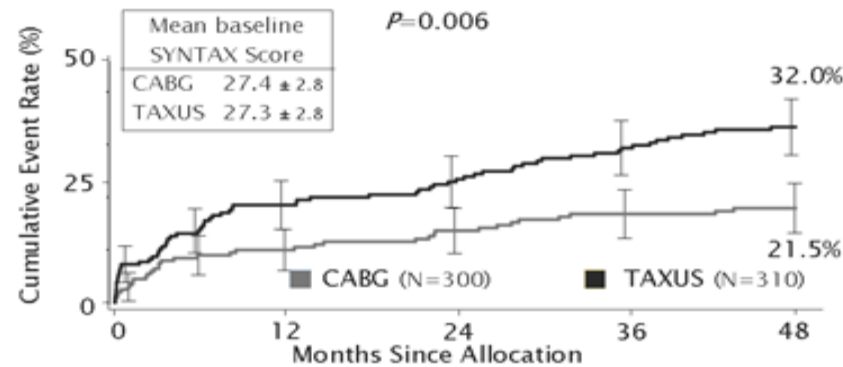
10. Heavy calcification

- a. ☐ No
b. ☐ Yes

11. Thrombus

- a. ☐ No
b. ☐ Yes

MACCE by SYNTAX Score 23-32



The cumulative MACCE rate is displayed for the SYNTAX Trial group this score corresponds to.

Summary

Lesion 1

(segment 1): 1x2=

Length >20 mm

Sub total lesion 1

2
1
3

Lesion 2

(segment 7): 2.5x2=

Bifurcation Type: Medina 1,1,1:

Length >20 mm

Heavy calcification

Sub total lesion 2

5
2
1
2
10

Lesion 3

segment number(s)

(segment 11): 1.5x5=

Age T.O. is yes

+ Blunt stump

the first segment beyond the T.O. visualized by contrast: 13

Heavy calcification

Sub total lesion 3

7.5
1
1
0
2
11.5

TOTAL:

24.5

What is Better

Web Calculator

Segments:	Lesions:	1
RCA	RCA proximal	1
	RCA mid	2
	RCA distal	3
	Posterior descending	4
	Posterolateral from RCA	16
	Posterolateral from RCA	16a
	Posterolateral from RCA	16b
	Posterolateral from RCA	16c
LM	Left main	5
LAD	LAD proximal	6
	LAD mid	7
	LAD apical	8
	First diagonal	9
	Add. first diagonal	9a
	Second diagonal	10
	Add. second diagonal	10a
LCX	Proximal circumflex	11
	Intermediate/anterolateral	12
	Obtuse marginal	12a
	Obtuse marginal	12b
	Distal circumflex	13
	Left posterolateral	14
	Left posterolateral	14a
	Left posterolateral	14b

4. Total occlusion (T.O.)

a. ☒ No
b. ☐ Yes:

5. Trifurcation

a. ☒ No
b. ☐ Yes

6. Bifurcation

Spread Sheet(Excel)

Database Program(ACCESS, mySQL)

Web Calculator

Spread Sheet(Excel)
Database Program(ACCESS, mySQL)

Input Limit



Serial	Hospital ID	Dominance	No. Lesion	Segment	Segment No	Seg Score	Total Occlusion	TO AGE	TO stump	TO Bridging	TO visual	TO SB	Trifur	Bifur	Angulation	Ostial Lesion	Tortuosity	Length 20>	Calcific	Thrombus	DDS1	DDS2	Lesion Score
B01R00001	12345678	0	5	1RCAPROXIMAL																			

- Segment Select

- ☒ 1. RCA PROXIMAL
- ☐ 2. RCA MID
- ☐ 3. RCA DISTAL
- ☐ 4. PDA[]
- ☐ 16. PL FROM RCA
- ☐ 16(a). PL FROM RCA
- ☐ 16(b). PL FROM RCA
- ☐ 16(c). PL FROM RCA
- ☐ 5. LEFT MAIN
- ☐ 6. LAD PROXIMAL
- ☐ 7. LAD MID
- ☐ 8. LAD APICAL
- ☐ 9. FIRST DIAGONAL
- ☐ 9(a). FIRST DIAGONAL(a)
- ☐ 10. SECOND DIAGONAL
- ☐ 10(a). SECOND DIAGONAL(a)
- ☐ 11. LCX PROXIMAL
- ☐ 12. INTERMEDIATE/ANTEROLATERAL ARTERY
- ☐ 12(a). OM(a)
- ☐ 12(b). OM(b)
- ☐ 13. LCX DISTAL
- ☐ 14. LEFT POSTEROLATERAL
- ☐ 14(a). LEFT POSTEROLATERAL
- ☐ 14(b). LEFT POSTEROLATERAL

- Trifurcation Select

- ☒ 0
- ☐ 10.0.0
- ☐ 0.10.0
- ☐ 0.0.10
- ☐ 0.0.0.1
- ☐ 110.0
- ☐ 10.10
- ☐ 10.0.1
- ☐ 0.110
- ☐ 0.10.1
- ☐ 0.0.11
- ☐ 1110
- ☐ 10.11
- ☐ 110.1
- ☐ 0.111

- Bifurcation Select

- ☐ 0
- ☐ 10.0
- ☐ 0.10
- ☐ 0.0.1
- ☐ 110
- ☐ 10.1
- ☐ 0.11
- ☒ 111

Input Select - Segment

	A	B	C	D	E	F	G	H	I	J	K
1	No.	Treatment_date	ID	DOMINAN (Right=0 Left=1)	No. LESION	SEGMENTS	Total Occlusion(TO)				
2							TO (0 ,1)	TO AGE >3MONT H or UNKNOWN (0 ,1)	BLUNT STUMP (0 ,1)	BRIDGING COLLATERALS (0 ,1)	FIRST SEGMENT VISIBLE BEYOND TO (0 ,1,2,3)
250	698	08-Jan-09	27613146	0	3	9. FIRST DIAGONAL	0	0	0	0	0
251	698	08-Jan-09	27613146	0	3	1. RCA PROXIMAL	0	0	0	0	0
252	699	08-Jan-09	12768943	0	4	1. RCA PROXIMAL	0	0	0	0	0
253	699	08-Jan-09	12768943	0	4	13. LCX DISTAL	0	0	0	0	0
254	699	08-Jan-09	12768943	0	4	6. LAD PROXIMAL	0	0	0	0	0
255	699	08-Jan-09	12768943	0	4	7. LAD MID	0	0	0	0	0
256	699	08-Jan-09	12768943	0	4	9. FIRST DIAGONAL	0	0	0	0	0
257	700	09-Jan-09	30762479	0	1						
258						1. RCA PROXIMAL					
259						2. RCA MID					
260						3. RCA DISTAL					
261						4. PDA					
262						16. PL FROM RCA					
263						16(a). PL FROM RCA					
						16(b). PL FROM RCA					
						16(c). PL FROM RCA					

Data Validation (데이터 유효성검사)

G258																			
1	A	B	C	D	E	F	Total Occlusion(TO)							N	BIFURCATION		Q	R	S
	No.	Treatment_date	ID	DOMINAN (Right=0 Left=1)	No. LESION	SEGMENTS	TO (0,1)	TO AGE >3MONTH H or UNKNOWN (0,1)	BLUNT STUMP (0,1)	BRIDGING COLLATERALS (0,1)	FIRST SEGMENT VISIBLE BEYOND TO (0,1,2,3)	SB<1.5 mm (0,1)	BOTH SB (0,1)	TRIFURCA TION MEDINA	BIFURCATI ON MEDINA	ANGULATIO N < 70° (0,1)	OS LESION (0,1)	BR OSTIUM LESION (0,1)	SEVERE TORTUOSITY (0,1)
2																			
250	698	08-Jan-09	27613146	0	3	9. FIRST DIAGONAL	0	0	0	0	0	0	0	0	0	0	0	0	0
251	698	08-Jan-09	27613146	0	3	1. RCA PROXIMAL	0	0	0	0	0	0	0	0	0	0	0	0	0
252	699	08-Jan-09	12768943	0	4	1. RCA PROXIMAL	0	0	0	0	0	0	0	0	0	0	0	0	0
253	699	08-Jan-09	12768943	0	4	13. LCX DISTAL	0	0	0	0	0	0	0	0	0	0	0	0	0
254	699	08-Jan-09	12768943	0	4	6. LAD PROXIMAL	0	0	0	0	0	0	0	0	0	0	0	0	0
255	699	08-Jan-09	12768943	0	4	7. LAD MID	0	0	0	0	0	0	0	0	0	0	0	0	0
256	699	08-Jan-09	12768943	0	4	9. FIRST DIAGONAL	0	0	0	0	0	0	0	0	0	0	0	0	0
257	700	09-Jan-09	30762479	0	1	2. RCA MID	0	0	0	0	0	0	0	0					
258																			
259																			
260																			
261																			
262																			
263																			
264																			
265																			
266																			
267																			
268																			
269																			
270																			
271																			

데이터 유효성

유효성 조건

제한 대상(A): 정수 ☒ 공백 무시(B)

제한 방법(D): 해당 범위

최소값(M): 0

최대값(X): 1

☐ 변경 내용을 설정이 같은 모든 셀에 적용(E)

모두 지우기(C) 확인 취소

데이터 유효성

유효성 조건

제한 대상(A): 정수 ☒ 공백 무시(B)

제한 방법(D): 해당 범위

최소값(M): 0

최대값(X): 3

☐ 변경 내용을 설정이 같은 모든 셀에 적용(E)

모두 지우기(C) 확인 취소

Data Validation (데이터 유효성검사)

1	A	B	C	D	E	F	Total Occlusion(TO)							N
	No.	Treatment_date	ID	DOMIN AN (Right=0 Left=1)	No. LESION	SEGMENTS	TO (0,1)	TO AGE >3MONT H or UNKNOWN	BLUNT STUMP (0,1)	BRIDGING COLLATERALS (0,1)	FIRST SEGME NT VISIBLE	SB<1.5 mm (0,1)	BOTH SB (0,1)	TRIFURCA TION MEDINA
239	696	07-Jan-09	18315998	0	6	12(a). OM(a)	0	0	0	0	0	0	0	0
240	696	07-Jan-09	18315998	0	6	7. LAD MID	0	0	0	0	0	0	0	0
241	696	07-Jan-09	18315998	0	6	6. LAD PROXIMAL	0	0	0	0	0	0	0	0
242	696	07-Jan-09	18315998	0	6	(a). FIRST DIAGONAL	0	0	0	0	0	0	0	0
243	696	07-Jan-09	18315998	0	6	2. RCA MID	0	0	0	0	0	0	0	0
244	697	08-Jan-09	28168416	0	5	13. LCX DISTAL	0	0	0	0	0	0	0	0
245	697	08-Jan-09	28168416	0	5	7. LAD MID	0	0	0	0	0	0	0	0
246	697	08-Jan-09	28168416	0	5	9. FIRST DIAGONAL	0	0	0	0	0	0	0	0
247	697	08-Jan-09	28168416	0	5	3. RCA DISTAL	0	0	0	0	0	0	0	0
248	697	08-Jan-09	28168416	0	5	6. LAD PROXIMAL	0	0	0	0	0	0	0	0
249	698	08-Jan-09	27613146	0	3	13. LCX DISTAL	0	0	0	0	0	0	0	0
250	698	08-Jan-09	27613146	0	3	9. FIRST DIAGONAL	0	0	0	0	0	0	0	0
251	698	08-Jan-09	27613146	0	3	1. RCA PROXIMAL	0	0	0	0	0	0	0	0
252	699	08-Jan-09	12768943	0	4	1. RCA PROXIMAL	0	0	0	0	0	0	0	0
253	699	08-Jan-09	12768943	0	4	13. LCX DISTAL	0	0	0	0	0	0	0	0
254	699	08-Jan-09	12768943	0	4	6. LAD PROXIMAL	0	0	0	0	0	0	0	0
255	699	08-Jan-09	12768943	0	4	7. LAD MID	0	0	0	0	0	0	0	0
256	699	08-Jan-09	12768943	0	4	9. FIRST DIAGONAL	0	0	0	0	0	0	0	0
257	700	09-Jan-09	30762479	0	1	2. RCA MID	0	0	0	0	0	0	0	0

데이터 유효성

설정 설명 메시지 오류 메시지 IME 모드

유효성 조건

제한 대상(S):

목록 ☐ 공백 무시(B) ☐ 드롭다운 표시(I)

제한 방법(D):

해당 범위

원본(S):

=BN\$3:BU\$3

☐ 변경 내용을 설정이 같은 모든 셀에 적용(P)

모두 지우기(C) 확인 취소

데이터 유효성

=BN\$3:BU\$3

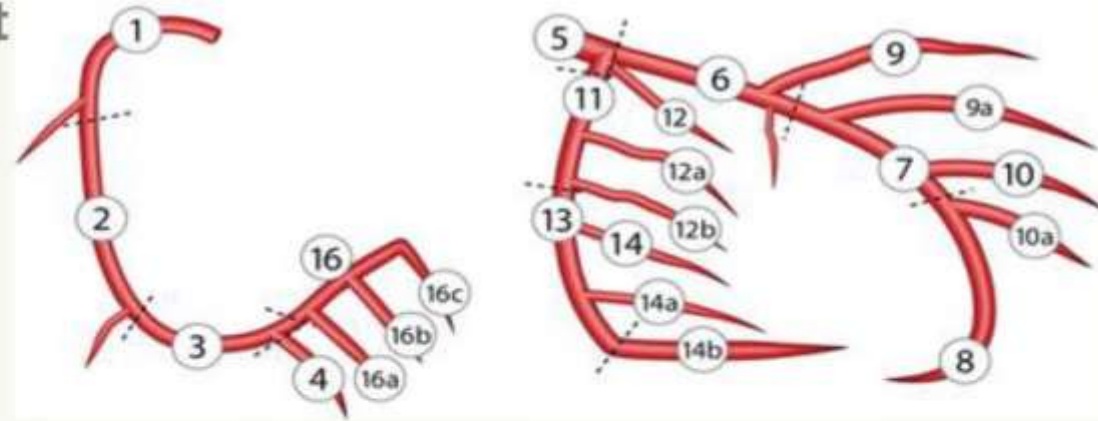
0 1.0.0 0.1.0 0.0.1 1.1.0 1.0.1 0.1.1 1.1.1

Segment Weighting Factors

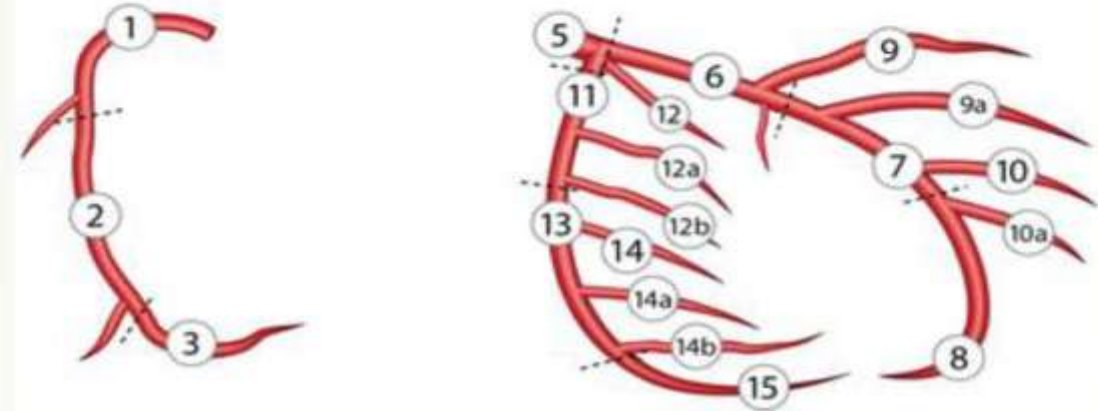
Table 1. Segment weighing factors

Segment No	Right dominance	Left dominance
1 RCA proximal	1	0
2 RCA mid	1	0
3 RCA distal	1	0
4 Posterior descending artery	1	n.a.
16 Posterolateral branch from RCA	0.5	n.a.
16a Posterolateral branch from RCA	0.5	n.a.
16b Posterolateral branch from RCA	0.5	n.a.
16c Posterolateral branch from RCA	0.5	n.a.
5 Left Main	5	6
6 LAD proximal	3.5	3.5
7 LAD mid	2.5	2.5
8 LAD apical	1	1
9 First diagonal	1	1
9a First diagonal ^a	1	1
10 Second diagonal	0.5	0.5
10a Second diagonal ^a	0.5	0.5
11 Proximal circumflex artery	1.5	2.5
12 Intermediate/ anterolateral artery	1	1
12a Obtuse marginal ^a	1	1
12b Obtuse marginal ^b	1	1
13 Distal circumflex artery	0.5	1.5
14 Left posterolateral	0.5	1
14a Left posterolateral ^a	0.5	1
14b Left posterolateral ^b	0.5	1
15 Posterior descending	n.a.	1

Right



Left



Segment Weighting Factors

Table 1. Segment weighing factors

Segment No	Right dominance	Left dominance
1 RCA proximal	1	0
2 RCA mid	1	0
3 RCA distal	1	0
4 Posterior descending artery	1	n.a.
16 Posterolateral branch from RCA	0.5	n.a.
16a Posterolateral branch from RCA	0.5	n.a.
16b Posterolateral branch from RCA	0.5	n.a.
16c Posterolateral branch from RCA	0.5	n.a.
5 Left Main	5	6
6 LAD proximal	3.5	3.5
7 LAD mid	2.5	2.5
8 LAD apical	1	1
9 First diagonal	1	1
9a First diagonal ^a	1	1
10 Second diagonal	0.5	0.5

Excel function (함수)

Dominance

=IF(AND(C2=0,G2=1),1,IF(AND(C2=0,G2=2),1,IF(AND(C2=0,G2=3),1,IF(AND(C2=0,G2=4),1,IF(AND(C2=0,G2=5),5,IF(AND(C2=0,G2=6),3.5,IF(AND(C2=0,G2=7),2.5,IF(AND(C2=0,G2=8),1,IF(AND(C2=0,G2=9),1,IF(AND(C2=0,G2=10),0.5,IF(AND(C2=0,G2=11),1.5,IF(AND(C2=0,G2=12),1,IF(AND(C2=0,G2=13),0.5,IF(AND(C2=0,G2=14),0.5,IF(AND(C2=0,G2=16),0.5,IF(AND(C2=1,G2=5),6,IF(AND(C2=1,G2=6),3.5,IF(AND(C2=1,G2=7),2.5,IF(AND(C2=1,G2=8),1,IF(AND(C2=1,G2=9),1,IF(AND(C2=1,G2=10),0.5,IF(AND(C2=1,G2=11),2.5,IF(AND(C2=1,G2=12),1,IF(AND(C2=1,G2=13),1.5,IF(AND(C2=1,G2=14),1,IF(AND(C2=1,G2=15),1,0)))))))))))))))))))))))))

14b	Left posterolateral ^b	0.5	1
15	Posterior descending	n.a.	1

Segment weight X multiplication factor



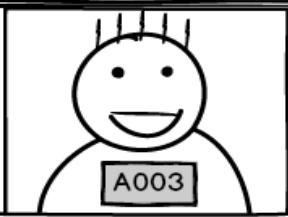
[illegible]

Dominance

=IF(AND(C2=0,G2=1),1,IF(AND(C2=0,G2=2),1,IF(AND(C2=0,G2=3),1,IF(AND(C2=0,G2=4),1,IF(AND(C2=0,G2=5),5,IF(AND(C2=0,G2=6),3.5,IF(AND(C2=0,G2=7),2.5,IF(AND(C2=0,G2=8),1,IF(AND(C2=0,G2=9),1,IF(AND(C2=0,G2=10),0.5,IF(AND(C2=0,G2=11),1.5,IF(AND(C2=0,G2=12),1,IF(AND(C2=0,G2=13),0.5,IF(AND(C2=0,G2=14),0.5,IF(AND(C2=0,G2=16),0.5,IF(AND(C2=1,G2=5),6,IF(AND(C2=1,G2=6),3.5,IF(AND(C2=1,G2=7),2.5,IF(AND(C2=1,G2=8),1,IF(AND(C2=1,G2=9),1,IF(AND(C2=1,G2=10),0.5,IF(AND(C2=1,G2=11),2.5,IF(AND(C2=1,G2=12),1,IF(AND(C2=1,G2=13),1.5,IF(AND(C2=1,G2=14),1,IF(AND(C2=1,G2=15),1,0)))))))))))))))))))))))))

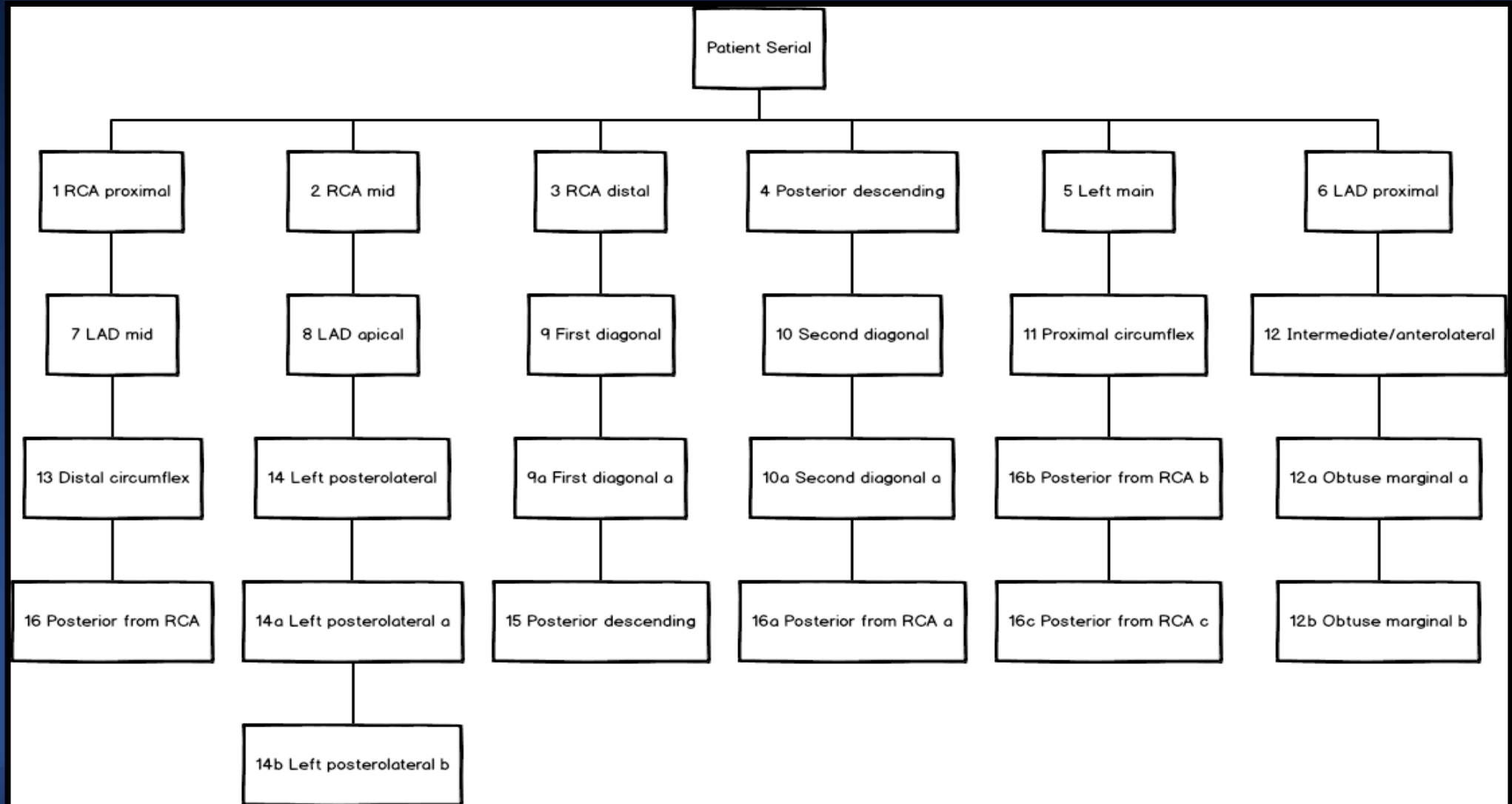
=(I2*J2)+SUM(K2:T2,V2:Y2)+COUNTA(Z2:AA2)

Lesion Score → Patient Score

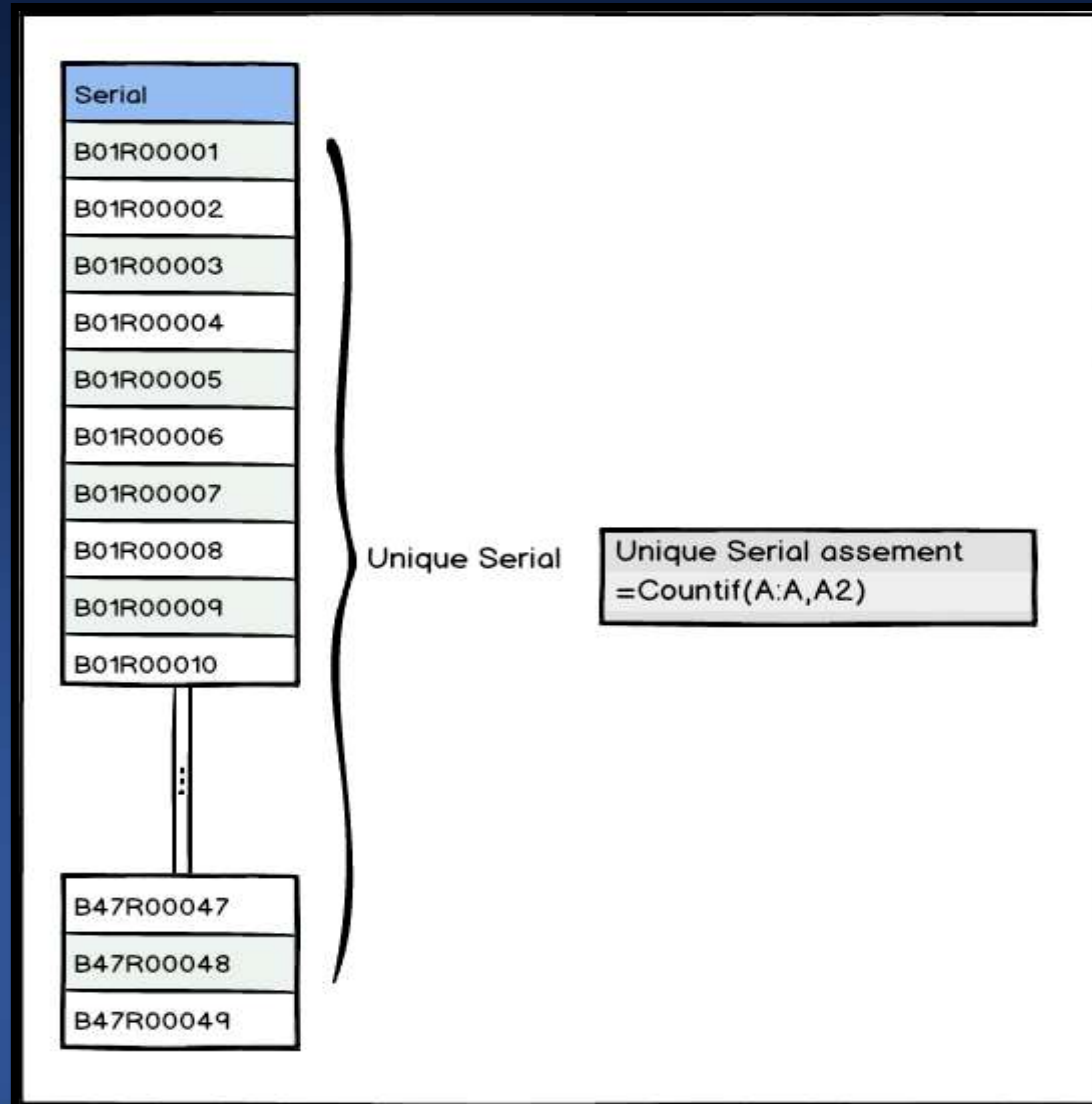
	Lesion No	Seg	Seg No	Lesion Score
	1	pRCA	Seg-1	5
	2	dRCA	Seg-3	3
	3	pLAD	Seg-6	8
	4	dLCX	Seg-11	2
	Lesion No	Seg	Seg No	Lesion Score
	1	mRCA	Seg-2	4
	2	Leftmain	Seg-5	11
	3	pLAD	Seg-6	10
	4	mLAD	Seg-7	6
	5	pLCX	Seg-11	7
	6	PL	Seg-16	1
	Lesion No	Seg	Seg No	Lesion Score
	1	mRCA	Seg-2	3
	2	pLAD	Seg-6	12
	3	pLCX	Seg-11	5

Patient	Seg-1	Seg-2	Seg-3	Seg-4	Seg-5	Seg-6	Seg-7	Seg-8	Seg-9	Seg-10	Seg-11	Seg-12	Seg-12a	Seg-13	Seg-14	Seg-15	Seg-16
A001	5		3			8					2						
A002		4			11	10	6				7						1
A003		3				12					5						

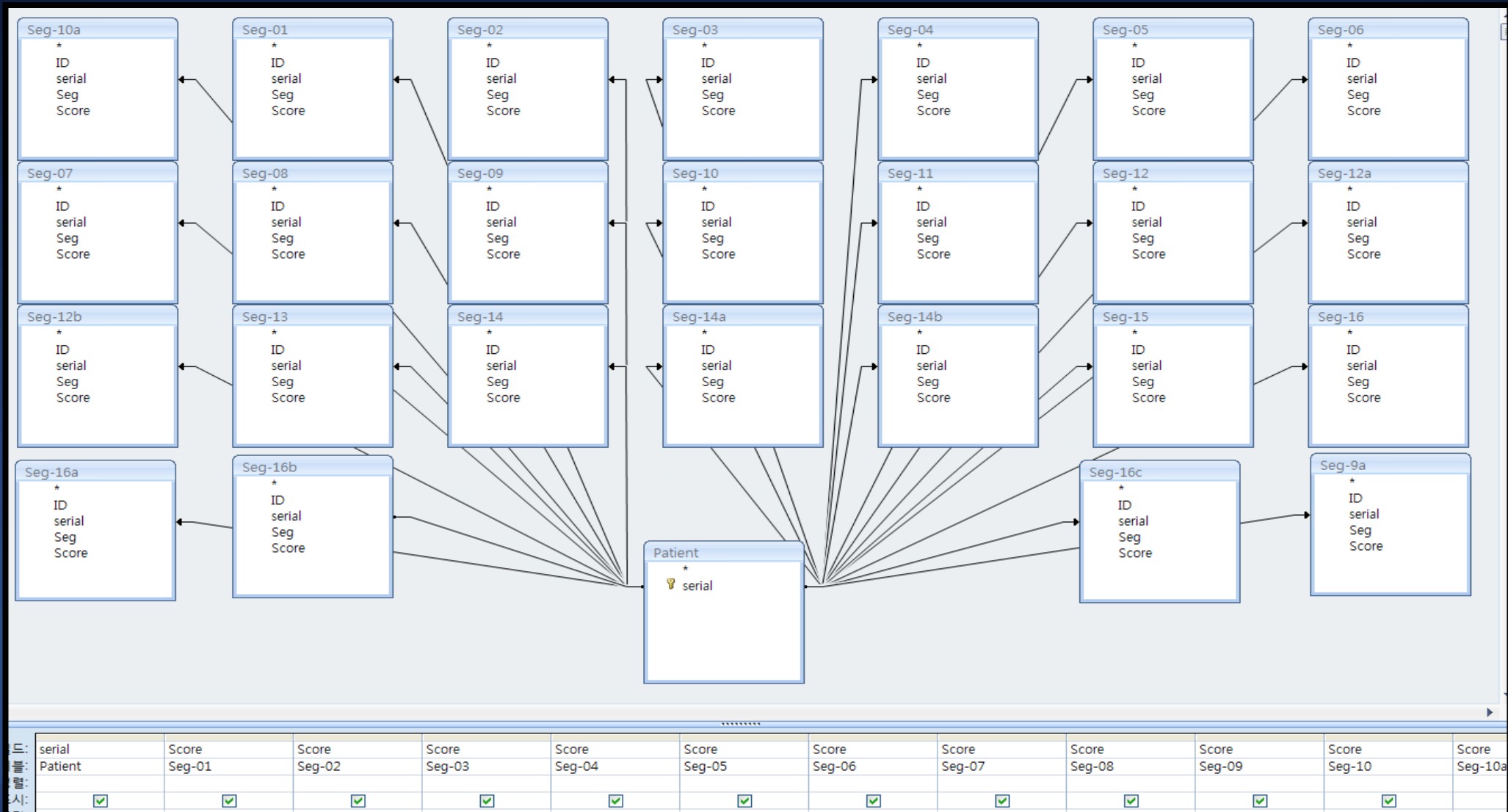
Segment Divide



Unique Serial



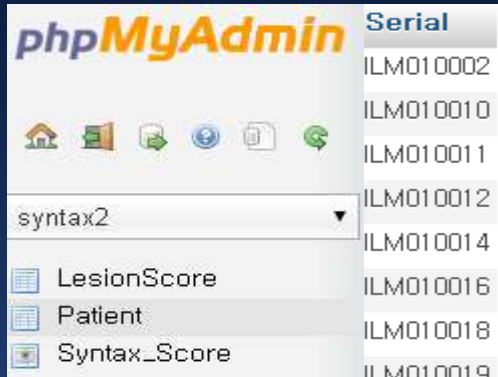
Join (Database Program : ACCESS)



Syntax Score

A	B	C	D	E	F	G	H	I	J	K	L	M	N	O	P	Q	R	S	T	U	V	W	X	Y	Z	AA
Serial	1	2	3	4	5	6	7	8	9	09a	10	10a	11	12	12a	12b	13	14	14a	14b	15	16	16a	16b	16c	Total
10748309		3	5		12		6	2					4				2									34
10767090	5	5			16	10	6																			42
11260136		2				8	6																			16
11939449					12	8							4													24
12094781			5	9	16		8																			38
12369416					13	8							3													24
12596845					12		13.5						6													31.5
12771057		4			11	7																				22
12802627					12		6										3									21
13166052		5			14	10																				29
13413347	4				15		5										4									28
13940957			3		14			2																		19
14086205					13																					13
15325686					10		12.5																			22.5
15427188		2			16									2												20
15691422		9			12		8						3													32
15750275					11	21.5																				32.5
15771986	2			3	12		9										2									28
15784595					14	22.5																				36.5
16261800	3	3			14																					20
16397730		3	6		17		8	3																		37
16441512						10											2									12
16656376					12			2									6.5									20.5

MySQL (Database Program)



Serial
ILM010002
ILM010010
ILM010011
ILM010012
ILM010014
ILM010016
ILM010018
ILM010019
ILM010020
ILM010021
ILM010022
ILM010024
ILM010025
ILM010026
ILM010028
ILM010030
ILM010033
ILM010038
ILM010040
ILM010041
ILM010042
ILM010043
ILM010048
ILM010049
ILM010051

Patient Serial



Serial	SEGMENT	Seg_Score
ILM010002	5	13
ILM010002	11	4
ILM010010	3	4
ILM010010	5	12
ILM010010	9	2
ILM010011	2	3
ILM010011	5	16
ILM010011	13	1
ILM010011	6	10
ILM010011	9	2
ILM010012	3	2
ILM010012	4	2
ILM010012	5	16
ILM010012	11	5
ILM010012	6	10
ILM010012	9	2
ILM010014	7	16.5
ILM010014	13	1
ILM010016	5	13
ILM010016	6	8
ILM010016	7	19.5
ILM010018	5	13
ILM010018	6	8
ILM010019	5	12

Lesion Score

SQL (Structure Query Language)

SELECT

```
`A`.`Serial` AS `Serial`,  
`SEG01`.`Seg_Score` AS `01_Score`,  
`SEG02`.`Seg_Score` AS `02_Score`,  
`SEG03`.`Seg_Score` AS `03_Score`,  
`SEG04`.`Seg_Score` AS `04_Score`,  
`SEG05`.`Seg_Score` AS `05_Score`,  
`SEG06`.`Seg_Score` AS `06_Score`,  
`SEG07`.`Seg_Score` AS `07_Score`,  
`SEG08`.`Seg_Score` AS `08_Score`,  
`SEG09`.`Seg_Score` AS `09_Score`,  
`SEG09a`.`Seg_Score` AS `09a_Score`,  
`SEG10`.`Seg_Score` AS `10_Score`,  
`SEG10a`.`Seg_Score` AS `10a_Score`,  
`SEG11`.`Seg_Score` AS `11_Score`,  
`SEG12`.`Seg_Score` AS `12_Score`,  
`SEG12a`.`Seg_Score` AS `12a_Score`,  
`SEG12b`.`Seg_Score` AS `12b_Score`,  
`SEG13`.`Seg_Score` AS `13_Score`,  
`SEG14`.`Seg_Score` AS `14_Score`,  
`SEG14a`.`Seg_Score` AS `14a_Score`,  
`SEG14b`.`Seg_Score` AS `14b_Score`,  
`SEG15`.`Seg_Score` AS `15_Score`,  
`SEG16`.`Seg_Score` AS `16_Score`,  
`SEG16a`.`Seg_Score` AS `16a_Score`,  
`SEG16b`.`Seg_Score` AS `16b_Score`,  
`SEG16c`.`Seg_Score` AS `16c_Score`
```

```
FROM (`Patient` `A` left join `LesionScore` `SEG01` on(`A`.`Serial` = `SEG01`.`Serial`) and (`SEG01`.`SEGMENT` = '1'))  
left join `LesionScore` `SEG02` on(`A`.`Serial` = `SEG02`.`Serial`) and (`SEG02`.`SEGMENT` = '2')  
left join `LesionScore` `SEG03` on(`A`.`Serial` = `SEG03`.`Serial`) and (`SEG03`.`SEGMENT` = '3')  
left join `LesionScore` `SEG04` on(`A`.`Serial` = `SEG04`.`Serial`) and (`SEG04`.`SEGMENT` = '4')  
left join `LesionScore` `SEG05` on(`A`.`Serial` = `SEG05`.`Serial`) and (`SEG05`.`SEGMENT` = '5')  
left join `LesionScore` `SEG06` on(`A`.`Serial` = `SEG06`.`Serial`) and (`SEG06`.`SEGMENT` = '6')  
left join `LesionScore` `SEG07` on(`A`.`Serial` = `SEG07`.`Serial`) and (`SEG07`.`SEGMENT` = '7')  
left join `LesionScore` `SEG08` on(`A`.`Serial` = `SEG08`.`Serial`) and (`SEG08`.`SEGMENT` = '8')  
left join `LesionScore` `SEG09` on(`A`.`Serial` = `SEG09`.`Serial`) and (`SEG09`.`SEGMENT` = '9')  
left join `LesionScore` `SEG09a` on(`A`.`Serial` = `SEG09a`.`Serial`) and (`SEG09a`.`SEGMENT` = '9a')  
left join `LesionScore` `SEG10` on(`A`.`Serial` = `SEG10`.`Serial`) and (`SEG10`.`SEGMENT` = '10')  
left join `LesionScore` `SEG10a` on(`A`.`Serial` = `SEG10a`.`Serial`) and (`SEG10a`.`SEGMENT` = '10a')  
left join `LesionScore` `SEG11` on(`A`.`Serial` = `SEG11`.`Serial`) and (`SEG11`.`SEGMENT` = '11')  
left join `LesionScore` `SEG12` on(`A`.`Serial` = `SEG12`.`Serial`) and (`SEG12`.`SEGMENT` = '12')  
left join `LesionScore` `SEG12a` on(`A`.`Serial` = `SEG12a`.`Serial`) and (`SEG12a`.`SEGMENT` = '12a')  
left join `LesionScore` `SEG12b` on(`A`.`Serial` = `SEG12b`.`Serial`) and (`SEG12b`.`SEGMENT` = '12b')  
left join `LesionScore` `SEG13` on(`A`.`Serial` = `SEG13`.`Serial`) and (`SEG13`.`SEGMENT` = '13')  
left join `LesionScore` `SEG14` on(`A`.`Serial` = `SEG14`.`Serial`) and (`SEG14`.`SEGMENT` = '14')  
left join `LesionScore` `SEG14a` on(`A`.`Serial` = `SEG14a`.`Serial`) and (`SEG14a`.`SEGMENT` = '14a')  
left join `LesionScore` `SEG14b` on(`A`.`Serial` = `SEG14b`.`Serial`) and (`SEG14b`.`SEGMENT` = '14b')  
left join `LesionScore` `SEG15` on(`A`.`Serial` = `SEG15`.`Serial`) and (`SEG15`.`SEGMENT` = '15')  
left join `LesionScore` `SEG16` on(`A`.`Serial` = `SEG16`.`Serial`) and (`SEG16`.`SEGMENT` = '16')  
left join `LesionScore` `SEG16a` on(`A`.`Serial` = `SEG16a`.`Serial`) and (`SEG16a`.`SEGMENT` = '16a')  
left join `LesionScore` `SEG16b` on(`A`.`Serial` = `SEG16b`.`Serial`) and (`SEG16b`.`SEGMENT` = '16b')  
left join `LesionScore` `SEG16c` on(`A`.`Serial` = `SEG16c`.`Serial`) and (`SEG16c`.`SEGMENT` = '16c'))  
order by `A`.`Serial`;
```

Syntax Score

Serial	01_Score	02_Score	03_Score	04_Score	05_Score	06_Score	07_Score	08_Score	09_Score	09a_Score	10_Score	10a_Score	11_Score	12_Score	12a_Score	12b_Score	13_Score	14_Score	14a_Score	14b_Score	15_Score	16_Score	16a_Score	16b_Score	16c_Score
UMD10002	NULL	NULL	NULL	NULL 13		NULL	NULL	NULL	NULL	NULL	NULL	NULL 4		NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL
UMD10010	NULL	NULL 4		NULL 12		NULL	NULL	NULL 2		NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL
UMD10011	NULL 3		NULL	NULL 15	10		NULL	NULL 2		NULL	NULL	NULL	NULL	NULL	NULL	NULL 1		NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL
UMD10012	NULL	NULL 2	2	16	10		NULL	NULL 2		NULL	NULL	NULL 5		NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL
UMD10014	NULL	NULL	NULL	NULL	NULL	NULL 15.5		NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL 1		NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL
UMD10016	NULL	NULL	NULL	NULL 13	8	19.5		NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL
UMD10018	NULL	NULL	NULL	NULL 13	8		NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL
UMD10019	NULL	NULL	NULL	NULL 12		NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL
UMD10020	NULL	NULL	NULL	NULL 14	13		NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL
UMD10021	NULL 3		NULL	NULL 12	10	6		NULL	NULL	NULL	NULL	NULL 11.5	2		NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL
UMD10022	NULL 11		NULL	NULL 15		NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL
UMD10024	NULL	NULL	NULL	NULL 10		NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL
UMD10025	NULL 6		NULL	NULL 11		NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL 1		NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL
UMD10026	NULL 3		NULL	NULL 12		NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL
UMD10028	NULL	NULL	NULL	NULL 12		NULL 6		NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL
UMD10030	NULL	NULL	NULL	NULL 13	11		NULL	NULL 3		NULL	NULL	NULL 7		NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL
UMD10033 5	5		NULL	NULL 20	12		NULL	NULL	NULL	NULL	NULL	NULL 4		NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL
UMD10038	NULL	NULL	NULL	NULL 14	8	6		NULL	NULL	NULL	NULL	NULL 5		NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL
UMD10040	NULL	NULL	NULL	NULL 10		NULL 15.5		NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL
UMD10041	NULL	NULL	NULL	NULL 12		NULL 6		NULL 2		NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL
UMD10042	NULL	NULL	NULL	NULL 12		NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL
UMD10043	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL
UMD10048	NULL	NULL	NULL	NULL 11		NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL 2		NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL
UMD10049	NULL	NULL	NULL	NULL 12	9		NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL
UMD10051	NULL	NULL	NULL	NULL 12	11		NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL
UMD10053 3		NULL 3		NULL 11		NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL	NULL

Conlcusion

- Spread sheet(Excel)을 통해 Syntax Score Data를 쉽게 입력
- Excel을 통해 입력을 할 경우 Data Validation (유효성 검사)기능으로 입력오류 예방
- Excel의 함수(Function)로 Segment weighting factors, Segment Score를 구할 수 있음
- Syntax Score(per patient)를 계산하기 위해 ACCESS, MySQL을 이용하면 효율적임