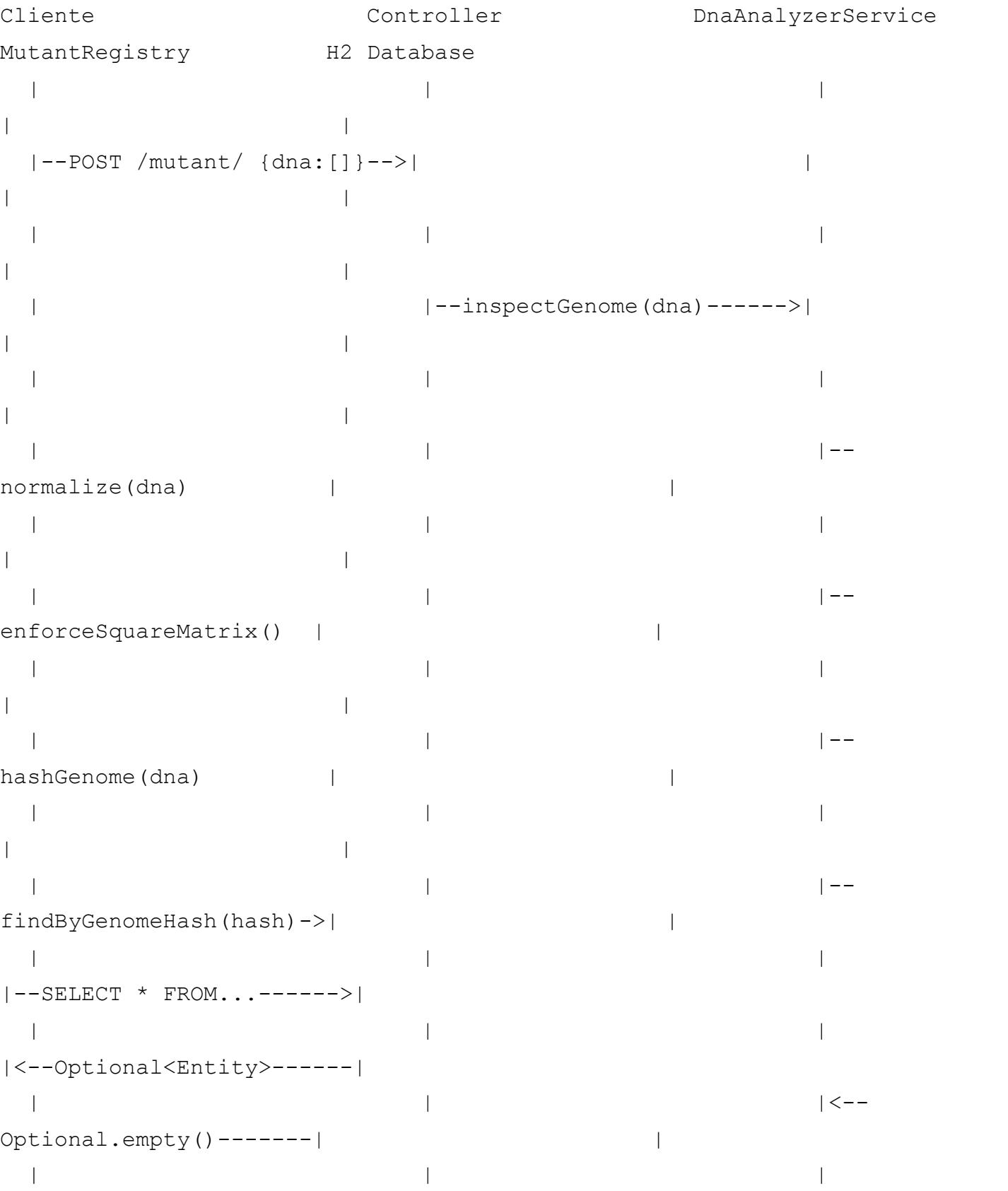


Diagrama de Secuencia - Mutant Detector API

Flujo: POST /mutant/



```

|
|
|
buildMatrix(dna)
|
|
|
|
|
detectMutations(matrix)
|
|
|
|
|
exploreRows()
|
|
|
exploreColumns()
|
|
|
explorePrimaryDiagonals()
|
|
|
exploreSecondaryDiagonals()
|
|
|
|
|
(mutant)-----|
|
|
|
|
persistResult()
|
|
|
|
|
|
|
|
|
save(entity)----->|
|
|--INSERT INTO...----->|
|
|<--Entity saved-----|
|
|<--true-----
-----|
|
|<--true (mutant)-----|
|
|<--200 OK-----|
|

```

Flujo: GET /stats

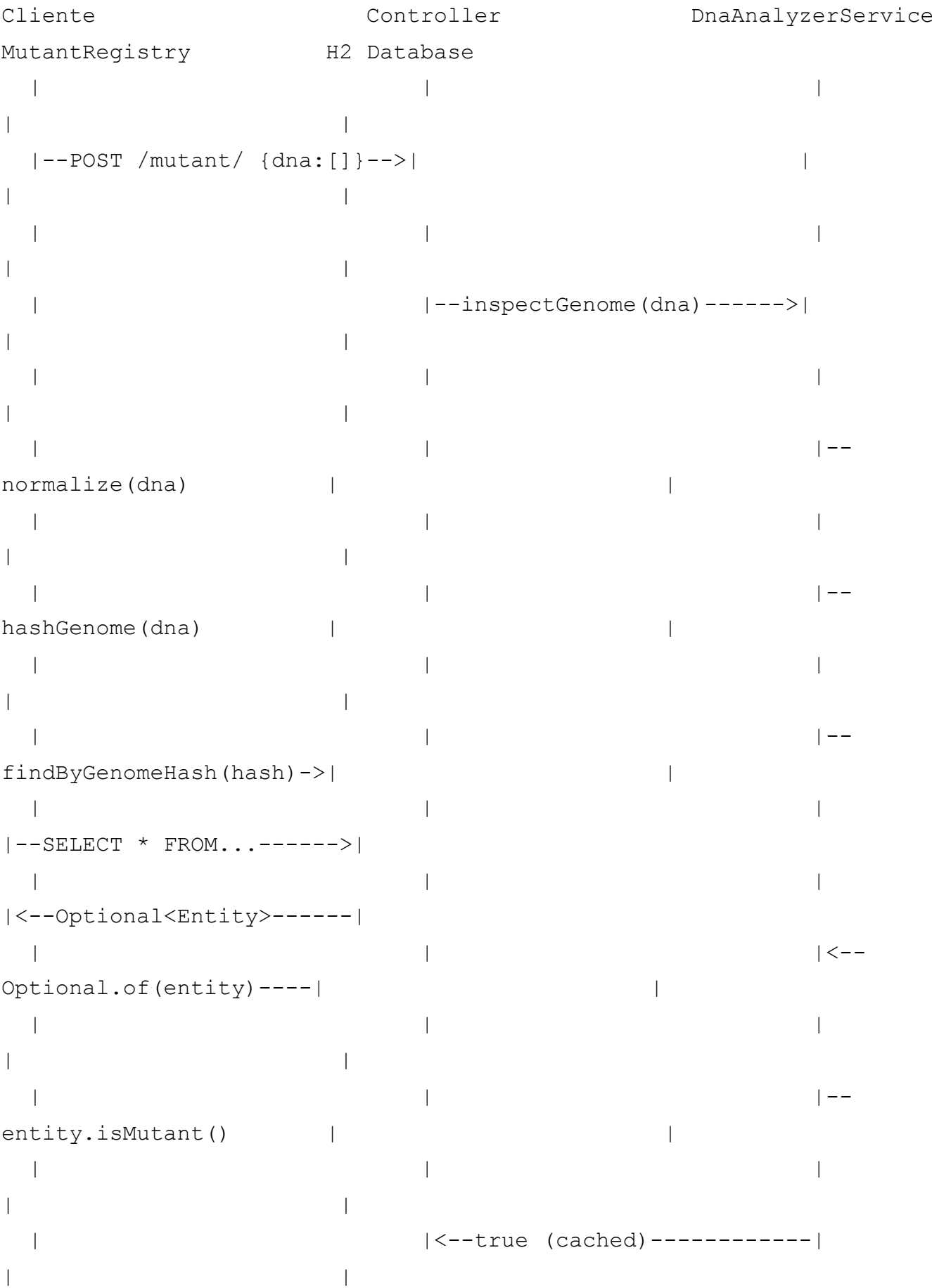
Cliente	Controller	DnaMetricsService
MutantRegistry	H2 Database	
--GET /stats----->		
	--fetchStats()----->	
		--
countByMutantTrue()----->		
--SELECT COUNT(*)...--->		
<--40-----		
		<--40-----

		--
countByMutantFalse()--->		
--SELECT COUNT(*)...--->		
<--100-----		
		<--100-----

		--calculate
ratio (40/100)		
		--new
StatsPayload(40, 100, 0.4)		
	<--StatsPayload-----	

```
|<--200 OK {count_mutant_dna:40, count_human_dna:100, ratio:0.4}--|
|
|
```

Flujo: POST /mutant/ (con caché)



|<--200 OK-----|

|

|

|