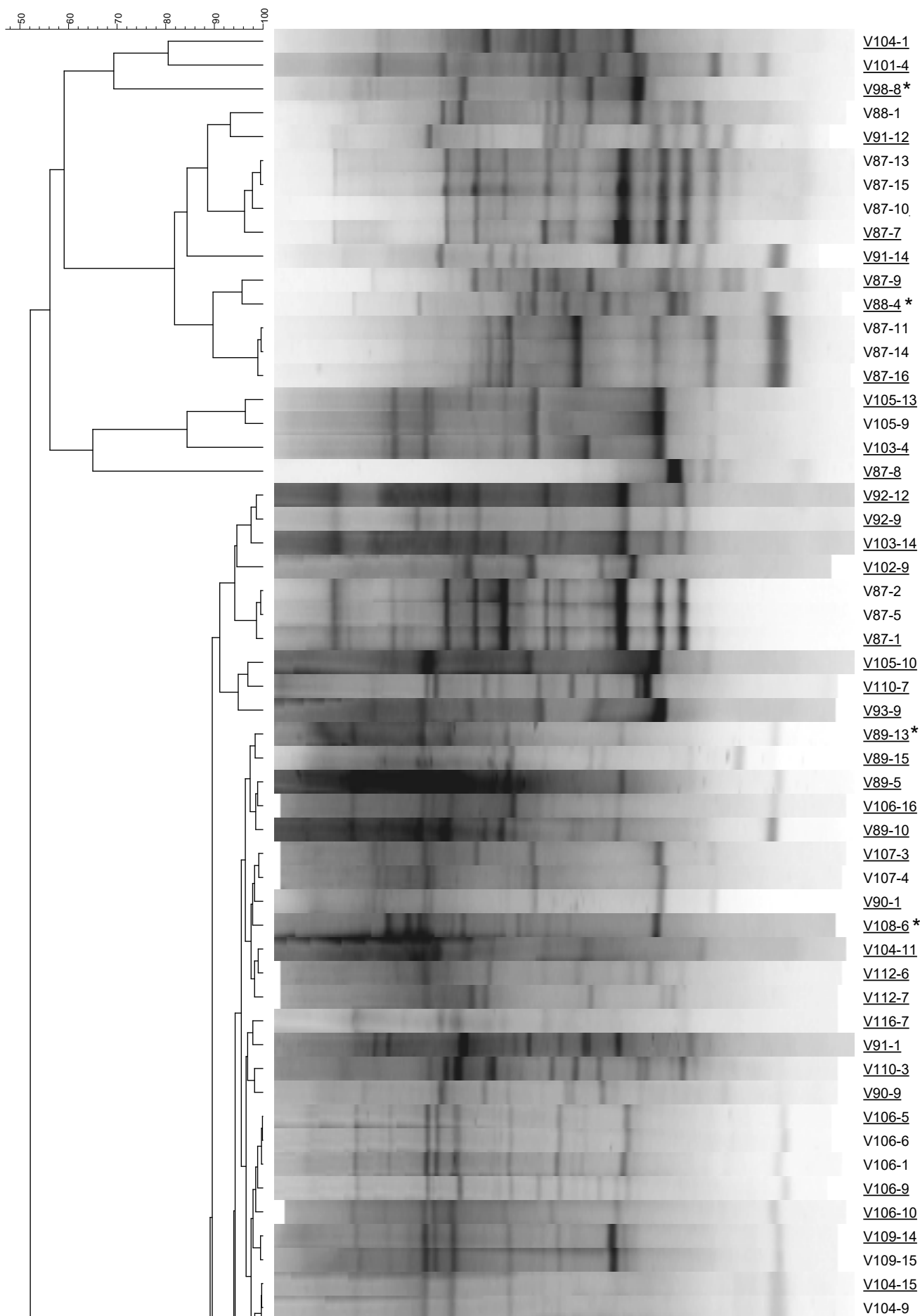
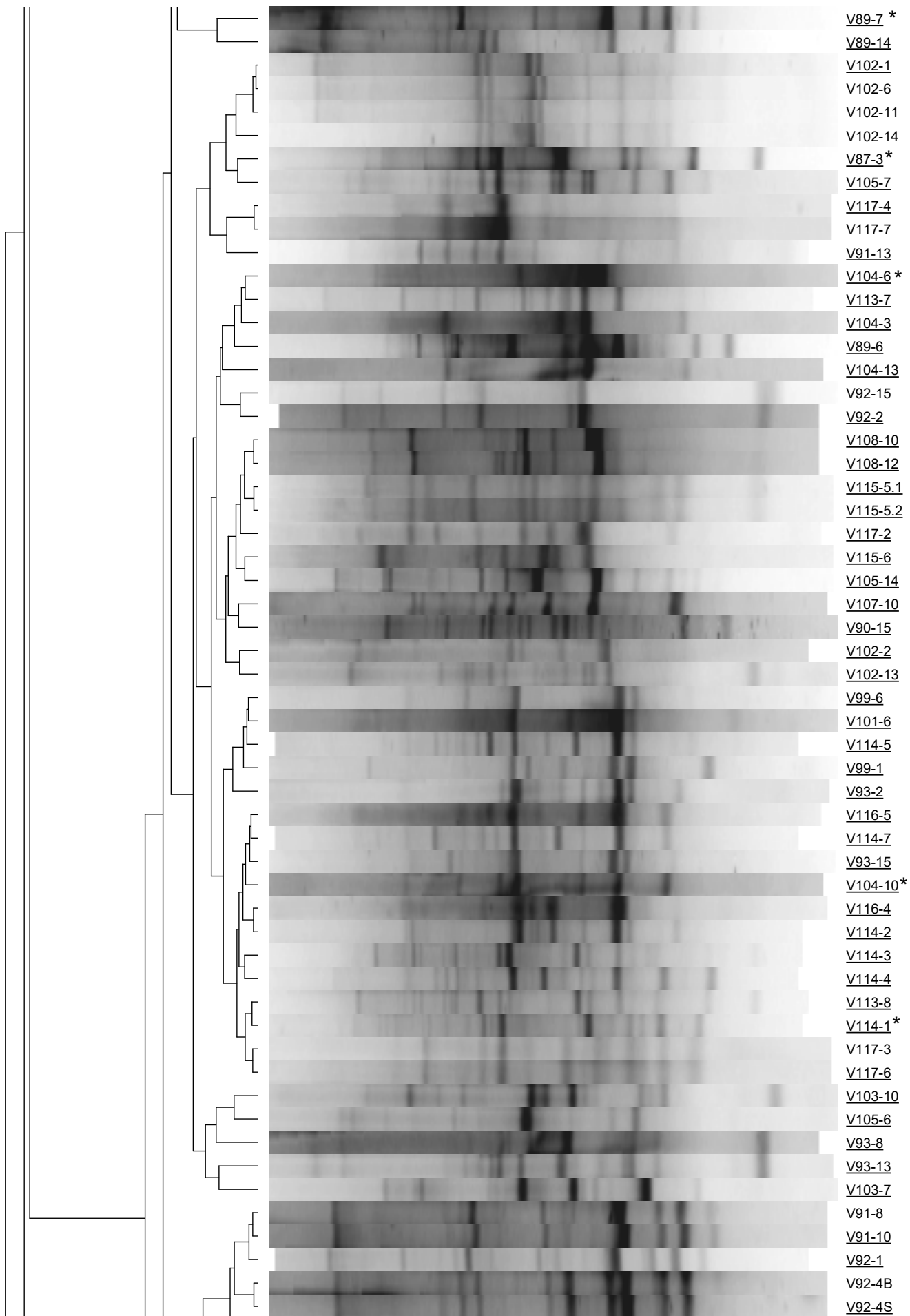


RAPD



	V104-14
	<u>V110-4</u>
	<u>V103-12</u>
	<u>V113-5</u>
	<u>V113-2</u>
	<u>V107-8</u>
	<u>V107-9</u>
	<u>V104-12</u>
	<u>V104-8</u>
	<u>V102-8</u>
	<u>V88-15</u>
	<u>V92-6</u>
	<u>V107-11</u>
	<u>V107-14</u>
	<u>V90-13</u>
	<u>V91-2</u>
	<u>V91-9</u>
	<u>V108-9</u>
	<u>V86-2</u>
	<u>V86-8</u>
	<u>V86-11</u>
	<u>V89-1</u>
	<u>V106-11</u> *
	<u>V108-7</u>
	<u>V90-5</u>
	<u>V106-3</u>
	<u>V108-4</u>
	<u>V93-10</u>
	<u>V91-6</u>
	<u>V93-11</u>
	<u>V93-14</u>
	<u>V87-6</u>
	<u>V106-4</u>
	<u>V103-11</u>
	<u>V103-1</u>
	<u>V103-15</u>
	<u>V86-16</u>
	<u>V103-8</u>
	<u>V86-14</u>
	<u>V90-4</u> *
	<u>V112-1</u>
	<u>V102-15</u>
	<u>V88-12</u>
	<u>V97-8</u>
	<u>V105-4</u>
	<u>V114-8</u>
	<u>V114-6</u>
	<u>V115-2</u>
	<u>V116-8.2</u>
	<u>V116-6.2</u>
	<u>V116-6.1</u>
	<u>V116-8.1</u>
	<u>V109-10</u>
	<u>V109-2</u>
	<u>V109-1</u>
	<u>V109-16</u>

		<u>V108-11</u>
		V108-15
		<u>V98-6</u>
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		<u>V91-4B</u>
		V91-4S
		<u>V98-7</u>
		<u>V109-7</u>
		V109-3
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		<u>V109-11</u>
		V86-13
		V86-15
		<u>V86-10</u>
		V86-6
		V86-3
		V86-4
		V86-1
		V86-9
		<u>V86-12</u>
		<u>V113-1</u>
		<u>V90-14 *</u>
		<u>V91-7</u>
		<u>V89-3</u>
		<u>V89-12</u>
		<u>V115-4</u>
		<u>V90-2</u>
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		<u>V90-16</u>
		<u>V89-11 *</u>
		<u>V104-4</u>
		<u>V89-4 *</u>
		V105-15
		<u>V105-16</u>
		<u>V115-8 *</u>
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		<u>V110-5</u>
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		<u>V106-13</u>
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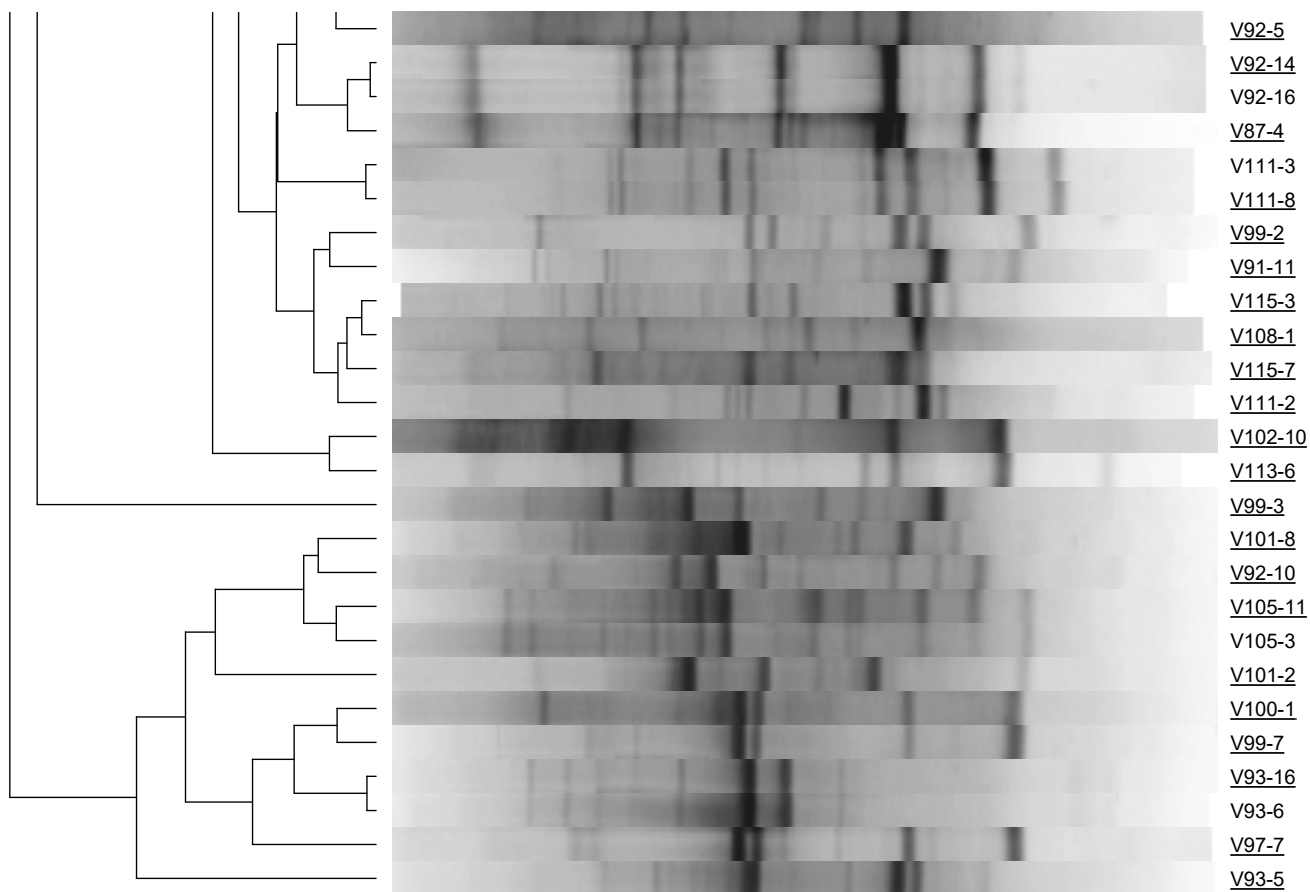


Figure S1: Dendrogram obtained by UPGMA clustering using the Pearson correlation coefficient of RAPD-PCR fingerprints of 248 Gram-negative isolates from lettuces and different fertilized soil samples. Selected strains for 16S rRNA gene sequencing were marked with an under bar. *; selected isolates for whole genome sequencing.